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| 4860487   | C 303  | 299<br>300<br>c 301<br>302   | c 296<br>c 296<br>. 297  | c 293<br>294                           | c 290<br>c 291<br>292   |  | c 285<br>c 286  | 283<br>284                               | 280<br>c 281   | c 277<br>278<br>279   | 275<br>276   |                                       | 269<br>c 270<br>c 271  | 267<br>268   | 200<br>200<br>4  | c 262<br>c 263   |   | C 258  |  | c 253  | 251<br>252   |  | c 247<br>c 248   | 245<br>245   | C 242  |
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| 0.77777   | 0.777  | 0.7<br>0.7<br>0.7  | 0.7  | 0.7                                    | 0.7<br>0.7<br>0.7   | 0.7                                      | 0.7   |  | 0.7  |   |  |                                       |  |  | 0.7  | 0.7  | 0.7   | 0.7  | 0.7  | 0.7  | 0.7  | 0.7  | 0.7  | 0.7  | 0.7  |
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| Sequence 229054,<br>Sequence 290348,<br>Sequence 401173,<br>Sequence 401174,<br>Sequence 419733,<br>Sequence 445052,<br>Sequence 445053,<br>Sequence 445053,<br>Sequence 445053,                              | equence 117447,<br>equence 126633,<br>equence 168700,<br>equence 205145,   | Sequence 699381,<br>Sequence 956983,<br>Sequence 115601,<br>Sequence 93243, A                                  | Sequence 14626, A<br>Sequence 59357, A<br>Sequence 151132,<br>Sequence 583946,               | Sequence 1160, Ap<br>Sequence 17, Appl | Sequence 15215, A Sequence 15215, A Sequence 12, Appl                                       | Sequence 175133,<br>Sequence 23, Appl    | Sequence 3130, Ap<br>Sequence 10, Appl                      | 38.                                      | 0 N H  | 11<br>39<br>45  | 16,  | 250                                   | Sequence 526622,<br>Sequence 567430,                                 | Sequence /38154,<br>Sequence 346667,<br>Sequence 500855,             | Sequence 166136,<br>Sequence 349668,                                 | Sequence 115595,<br>Sequence 115596,                     | 0693, A   | Sequence 33, Appl<br>Sequence 98, Appl                         | equence 14, Ap   | 33160,   | nce 37171,<br>nce 648, <i>P</i>                                | nce 6145,  | nce 1800,  | Sequence 63/3, App<br>Sequence 823, App<br>Sequence 1311, Ap | nce 3080,  |
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9 US-10-325-899-5027
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Sequence 954464,
Sequence 118718,
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                                                                           RESULT 1

US-09-749-280B-127

Sequence 127, Application US/09749280B

GENERAL INFORMATION:

APPLICANT: Shimkets, Richard A.

APPLICANT: Leach, Martin

TITLE OF INVENTION: Nucleotide Polymorphisms and FILE REFERENCE: 15966-610

CURRENT APPLICATION NUMBER: US/09/749,280B

CURRENT FILING DATE: 2000-12-27

PRIOR APPLICATION NUMBER: 60/173,419

PRIOR PILING DATE: 1999-12-28

NUMBER OF SEQ ID NOS: 10038

SOFTWARE: CuraGen Patent Formatter Version 0.9

SEQ ID NO 127

LENGTH: 51

TYPE: DNA

ORGANISM: Homo sapiens

FEATURE:

NAME/KEY: allele
LOCATION: (26)...(0)
: LOCATION: (26)...(0)
: OTHER INFORMATION: Single nucle
: FEATURE:
: NAME/KEY: misc_feature
: LOCATION: (0)...(0)
: OTHER INFORMATION: Accession nu
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US-60-427-836-319951
US-60-427-836-306970
US-60-427-836-306970
US-60-427-836-301382
US-60-427-836-4297636-446242
US-60-427-836-4297836-446242
US-60-427-836-49917
1 US-60-427-836-49917
1 US-60-427-836-589418
US-09-94-1968-6
US-10-277-216-253
US-09-904-1968-6
US-10-318-966-6
US-10-318-966-6
US-10-399-73-239
US-09-992-633-239
US-09-992-633-378-399-143-632
US-10-349-143-638
1 US-10-35-899-148-9
1 US-10-35-899-149-9
1 US-10-325-899-1404
9 US-10-325-899-1404
              number cg44031583
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 509
LENGTH: 41
TYPE: DNA
ORGANISM: Human
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                                                                     Matches
                                                                                     Query Match
Best Local :
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Matches 3
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CURRENT FILING DATE: 2003-02-20
                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/09/749,280B CURRENT FILING DATE: 2000-12-27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION: Nucleic Acids Containing S TITLE OF INVENTION: Nucleotide Polymorphisms FILE REFERENCE: 15966-610
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Shimkets, Richard A. APPLICANT: Leach, Martin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: NUCLEOTIDE AND AMINO ACID SEQUENCES TITLE OF INVENTION: RELATING TO RESPIRATORY DISEASES A
                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: 60/173,419 PRIOR FILING DATE: 1999-12-28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILE REFERENCE: HUM02-11P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Keith, Tim
                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQ ID NOS: 10038
                                                                                                                                                     NAME/KEY: misc_feature
LOCATION: (0)...(0)
OTHER INFORMATION: Accession
                                                                                                                                                                                                               NAME/KEY: allele
LOCATION: (26)...(0)
OTHER INFORMATION: single nucleotide polymorphism
                                                                                                                                                                                                                                                                                                                 TYPE: DNA
                                                                                                                                                                                                                                                                                               ORGANISM: Homo sapiens
                                                                                                                                                                                                             FEATURE
                                                                                                                                                                                                                                                                              FEATURE:
                                                                                                                                                                                                                                                                                                                                 ENGTH: 51
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les 36; Conser
                                                                   Local Similarity
les 37; Conserv
                             1412 AAGGACACACAGGAGAGAGCCTTATGTATGCACAGAATGTGGG 1457
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     213 GTGGCATTCAGGGATGTGGCTGTGTACTTCACCCAGGAGGA 253
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39; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GTGACATTCAGGGATGTGGCCGTGTTCTTCAGCCAGGACGA 41
ACGGATACACACGGGAGAGAGCCTTATCAGTGCAAGGAGTGTGGG
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87.8%;
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Pred. No. 32;
0; Mismatches
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Pred. No. 44;
0; Mismatches
                                                                 Score 31.6; DB 5;
Pred. No. 1.2e+02;
0; Mismatches 9;
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RESULT 5 PCT-US03-02358-36

Sequence 36, Application PC/TUS0302358 GENERAL INFORMATION:

APPLICANT: Sera,

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RESUL: *
US-09-912-293-242606/c
Commence 242606, Application US/09912293
                                                                                                                                       , OTHER INFORMATION: n is equal to US\hbox{-}09\hbox{-}912\hbox{-}293\hbox{-}24\,2606
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                                                                                   Query Match
Best Local Similarity
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PRIOR APPLICATION NUMBER: 08/103,744
PRIOR FILING DATE: 1993-08-09
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Remaining Prior Application data removed - See NUMBER OF SEQ ID NOS: 244538
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION NUMBER: 09/813,155 PRIOR FILING DATE: 2001-03-21
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CURRENT FILING DATE: 2001-07-26
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FILING DATE: 1994-03-31
APPLICATION NUMBER: 09/741,830
FILING DATE: 2000-12-22
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                    2458 CTGAATTGAGAACCTGTCTTCCCATTTGGTGTGTCTTCCTCCGATTGATCCCAACC 2513
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE: 1994-03-3
55 CTGGATTGAGANCCTGTCTTTCCNTTTGG-CTGCTTTTCTNTGATTGATCNTCNCC
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                                                                      Conservative
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US-09-911-261A-36
; Sequence 36, Application US/09911261A
; GENERAL INFORMATION:
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; LCCATION: (45)..(56)
; OTHER INFORMATION: Nucleotides
US-09-911-2618-36
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US-09-911-261B-36
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQ ID NO 36
LENGTH: 60
TYPE: DNA
                                                                                                                                                                                                                                                                                              Query Match
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PRIOR FILING DATE: 2003-01-23
PRIOR FILING DATE: 2002-01-23
NUMBER OF SEQ ID NOS: 87
SOFTWARE: Patentin version 3.2
SEQ ID NO 36
                                                APPLICANT: Sera, Takashi
TITLE OF INVENTION: Zinc Finger Domain Recognition
FILE REFERENCE: 109845.135
                                                                                                                                                                                                                                                                 Matches
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           CURRENT APPLICATION NUMBER: US/09/911,261A CURRENT FILING DATE: 2001-07-23 PRIOR APPLICATION NUMBER: US 60/220,060
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                                                                                                                                                                                                                                                                                                                                                                                                                                  ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                        FEATURE:
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                                                                                                                                                                                                                                                             Local Similarity 60.0 es 36; Conservative
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Pred. No.
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Best Local
TITLE OF INVENTION: Nucleic Acids Containing Single TITLE OF INVENTION: Nucleotide Polymorphisms and Mc FILE REFERENCE: 15966-610 CURRENT APPLICATION NUMBER: US/09/749,280B CURRENT FILING DATE: 2000-12-27
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US-09-749-280B-6407
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Best Local Similarity
Marches 36; Conserv
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                                 Sequence 6407, Application US/09749280B GENERAL INFORMATION:
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NUMBER OF SEQ ID NOS: 69
SOFTWARE: PatentIn version 3.0
SEQ ID NO 36
APPLICANT: Shimkets, Richard APPLICANT: Leach, Martin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION NUMBER: 60/173,419
PRIOR FILING DATE: 1999-12-28
NUMBER OF SEQ ID NOS: 10038
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APPLICANT: Leach, Martin
TITLE OF INVENTION: Nucleic Acids Containing Single
TITLE OF INVENTION: Nucleotide Polymorphisms and M
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: CuraGen Patent Formatter Version
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILE REFERENCE: 15966-610
                                                                                                                                                                                                                                                                                                                   NAME/KEY: misc_feature LOCATION: (0)...(0)
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                                                                                                                                                                                                                                                                                                                                                                                                         NAME/KEY: allele
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TYPE: DNA
ORGANISM: Artificial Sequence
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                                                                                                                                                                                                          Local Similarity 73.9 nes 34; Conservative
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Pred. No. 3.1e+03;
0; Mismatches 12
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Pred. No.
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PCT-US03-02358-49
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US-09-911-261B-49
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PCT-US03-02358-49
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LENGTH: 56
TYPE: DNA
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Best Local Similarity
Matches 35; Conserv
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LENGTH: 51
           SEQ ID NO 49
LENGTH: 56
                                                                                                             APPLICANT: Sera, Takashi
TITLE OF INVENTION: Zinc Finger Domain Recognition Code and
FILE REFERENCE: 10845.135
CURRENT APPLICATION NUMBER: US/09/911,261B
CURRENT FILING DATE: 2000-07-21
CURRENT FILING DATE: 2000-07-26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION NUMBER: US 10/057,408
PRIOR FILING DATE: 2002-01-23
NUMBER OF SEQ ID NOS: 87
SOFTWARE: Patentin version 3.2
                                               PRIOR APPLICATION NUMBER: US 60/220,060
PRIOR FILING DATE: 2000-07-21
NUMBER OF SEQ ID NOS: 69
SOFTWARE: Patentin version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION NUMBER: PCT/US03/02358 CURRENT FILING DATE: 2003-01-23
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TITLE OF INVENTION: Zinc Finger Domain Recognition
FILE REFERENCE: 109845.158
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COTHER INFORMATION: Accession number cg43051067
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LOCATION: (26)...(0)
OTHER INFORMATION: single nucleotide polymorphism
TYPE: DNA
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ORGANISM: Homo sapiens
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nilarity 71.4%;
Conservative
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US-09-911-261A-49
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                                                                                                                                                                                                                 SEQ ID NO 45
LENGTH: 60
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Best Local :
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                                                                 Matches
                                                                                                Query Match
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TITLE OF INVENTION: Zinc Finger Domain Recognition Code and Uses Thereof
FILE REFERENCE: 109845.158
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TITLE OF INVENTION:
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                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: PCT/US03/02358 CURRENT FILING DATE: 2003-01-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             IIILE OF INVENTION: Zinc Finger Domain Recognition Code and Uses Thereof FILE REFERENCE: 109845.135
                                                                                                                                                                                                                                                  SOFTWARE: PatentIn version 3.2
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ORGANISM: Artificial Sequence
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GGGGAGAAGCCGTATAAATGTCCGGAATGTGGTAAAAGTTTTAGC
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Pred. No. 3.7e
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                                                                                Score 25.8; DB 1; Pred. No. 6.4e+03;
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Best Local Similarity
Watches 33; Conserve
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Sequence 45, Application US/09911261A
GENERAL INFORMATION:
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                                                                             Sequence 40, Application PC/TUS0302358
GENERAL INFORMATION:
APPLICANT: Sera, Takashi
TITLE OF INVENTION: Zinc Finger Domain Recognition Code and Uses
FILE REFERENCE: 109845.158
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LENGTH: 60
 PRIOR APPLICATION NUMBER: US 10/057,408
PRIOR FILING DATE: 2002-01-23
NUMBER OF CT.
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APPLICANT: Sera, Takashi
                                              CURRENT APPLICATION NUMBER: PCT/US03/02358
CURRENT FILING DATE: 2003-01-23
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NUMBER OF SEQ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Sera, Takashi
TITLE OF INVENTION: Zinc Finger Domain Recognition Code and Uses Thereof
FILE REFERENCE: 109845.135
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FILE REFERENCE: 109845.135
CURRENT APPLICATION NUMBER: US/09/911,261B
CURRENT FILING DATE: 2000-07-21
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TYPE: DNA
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33; Conserv
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Pred. No. 6.
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6.4e+03;
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US-09-911-261A-40
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US-09-911-261B-40
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                               SEQ ID NO 40
LENGTH: 56
                                                                                                                                                                                                           Sequence 40, Application US/09911261A GENERAL INFORMATION:
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Best Local
                                                                           PRIOR APPLICATION NUMBER: US 60/220,060 PRIOR FILING DATE: 2000-07-21 NUMBER OF SEQ ID NOS: 69
                                                                                                                              CURRENT APPLICATION NUMBER: US/09/911,261A CURRENT FILING DATE: 2001-07-23
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                                                               SOFTWARE: PatentIn version 3.0
                                                                                                                                                             APPLICANT: Sera, Takashi
TITLE OF INVENTION: Zinc Finger Domain Recognition Code and Uses Thereof
FILE REFERENCE: 109845.135
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TITLE OF INVENTION: Zinc Finger Domain Recognition Code and Uses Thereof.
ORGANISM: Artificial Sequence
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                  TYPE: DNA
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TYPE: DNA
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Pred. No. 7
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RESULT 20
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US-10-325-899-2511
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US-10-325-899-2511
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Best Local S
Matches 31
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Best Local
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LENGTH: 50
                                                                                                                                                                                                                                                                                             Sequence 3660, Application US/10325899 GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILE REFERENCE: 506612000122
CURRENT APPLICATION NUMBER: US/10/325,899
CURRENT FILING DATE: 2002-12-20
                                PRIOR APPLICATION NUMBER: US 10/006,290 PRIOR FILING DATE: 2001-10-22 PRIOR APPLICATION NUMBER: US 10/131,831
                                                                                      CURRENT APPLICATION NUMBER: US/10/325,899
CURRENT FILING DATE: 2002-12-20
PRIOR APPLICATION NUMBER: US 60/296,764
PRIOR FILING DATE: 2001-06-08
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: US 10/131,831 PRIOR FILING DATE: 2002-04-24
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Wohlgemuth,
                                                                                                                                                                                TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR DIAGNOSING AND MONITORING TRANSPLANTITLE OF INVENTION: REJECTION
                                                                                                                                                                                                                                                           APPLICANT:
                                                                                                                                                                                                                                                                          APPLICANT: Wohlgemuth, Jay
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 NUMBER OF
                                                                                                                                                                 FILE REFERENCE: 506612000122
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                   PRIOR FILING DATE:
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LOCATION: (48)..(58)
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mes 31; Conser
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33; Conser
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 SEQ ID NOS: 9966
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                                                                                                                                                                                                                                                                                                                                                                                                         CTTCAGTCAGAGCTCAAACCTTAGTCAACACCAGAGAATTCACATG 49
                                                                                                                                                                                                                   Fry, Kirk
Ly, Ngoc
Woodward, Robert
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Pred. No. 7
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Pred. No. 8.9e+03;
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7.1e+03;
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OTHER INFORMATION: Partial zinc finger domain oligomer
                   FEATURE:
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                                                                     ENGTH:
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; TYPE: DNA; Homo sapiens; ORGANISM: Homo sapiens US-10-325-899-3660
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Best Local
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                                                          SOFTWARE: PatentIn version 3.2 SEQ ID NO 50
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Best Local Similarity
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                                                                                                                                                              TITLE OF INVENTION: Zinc Finger Domain Recore FILE REFERENCE: 109845.158
CURRENT APPLICATION NUMBER: PCT/US03/02358
CURRENT FILING DATE: 2003-01-23
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                                                                                                      NUMBER OF SEQ ID NOS:
                                                                                                                         PRIOR APPLICATION NUMBER: US 10/057,408 PRIOR FILING DATE: 2002-01-23
                                                                                                                                                                                                                                                   APPLICANT: Sera,
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LOCATION: (0)...(0)
OTHER INFORMATION: Accession number cg43051067
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OTHER INFORMATION: single nucleotide polymorphism
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                     TYPE: DNA
ORGANISM: Artificial Sequence
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33; Conserv
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Pred. No. 8.9e+03;
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RESULT 24
US-09-911-261A-50/c
; Sequence 50, Application US/09911261A
; GENERAL INFORMATION:
RESULT 25
US-09-749-280B-5867
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                                                                                                                                                                                        ; OTHER INFORMATION: Partial zinc finger domain oligomer US-09-911-261a-50 \,
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                                                                                                                                                                                                                                                                    SEQ ID NO 50
LENGTH: 55
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                                                                                                                                            Query Match
Best Local
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APPLICANT: Sera, Ta
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                                                                                                                                                                                                                                                                                              NUMBER OF SEQ ID NOS: 69
SOFTWARE: PatentIn version 3.0
                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION NUMBER: US/09/911,261A CURRENT FILING DATE: 2001-07-23 PRIOR APPLICATION NUMBER: US 60/220,060 PRIOR FILING DATE: 2000-07-21
                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Sera, Takashi
TITLE OF INVENTION: Zinc Finger Domain Recognition Code and Uses Thereof
FILE REFERENCE: 109845.135
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION: Zinc Finger Domain Recognition Code and Uses Thereof FILE REFERENCE: 109845.135
CURRENT APPLICATION NUMBER: US/09/911,261B
CURRENT FILING DATE: 2000-07-21
PRIOR APPLICATION NUMBER: US 60/220,060
PRIOR FILING DATE: 2000-07-21
PRIOR FILING DATE: 2000-07-21
PRIOR FILING DATE: 2000-07-21
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                                                                                                                                                                                                                                                        TYPE: DNA
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                                                                                                                          Local Similarity
les 36; Conserv
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les 36; Consert
                                                                                 1211 CTTTAGCCTGAAGTCAAACCTCATTACCCACCAGAGGGGGGGCACACTGGGGGAGAAG 1265
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Pred. No. 1.
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Pred. No. 1
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Pred. No. 1.4e+04;
                                                                                                                     Mismatches
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l.4e+04;
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                                                                            Matches
                                                                                                              Query Match
                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: CuraGen Patent Formatter Version 0.9 SEQ ID NO 6408
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
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Best Local Similarity
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SOFTWARE: CuraGen Patent Formatter Version
SEQ ID NO 5867
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Shimkets, Richard A.
APPLICANT: Leach, Martin
TITLE OF INVENTION: Nucleic Acids Containing Single
TITLE OF INVENTION: Nucleotide Polymorphisms and Methods of Use Thereof
FILE REFERENCE: 15966-610
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CURRENT FILING DATE: 2000-12-27
PRIOR APPLICATION NUMBER: 60/173,419
PRIOR FILING DATE: 1999-12-28
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                                                                                                                                                                                                                                   NAME/KEY: allele LOCATION: (26)...(COTHER INFORMATION:
                                                                                                                                                          OTHER INFORMATION: Accession number cg43051067
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION: Nucleic Acids Containing Single TITLE OF INVENTION: Nucleotide Polymorphisms and Methods of Use Thereof FILE REFERENCE: 15966-610
                                                                                                                                                                                     NAME/KEY: misc_feature LOCATION: (0)...(0)
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                                                                                                                                                                                                                                                                                                                                   TYPE: DNA
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LOCATION: (26)...(0)
OTHER INFORMATION: single nucleotide polymorphism
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                                                                                             Similarity
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Homo sapiens
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Pred. No. 1.5e+04;
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Pred. No. 1
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US-09-911-261B-41/c
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RESULT 29
US-09-911-261A-41/c
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Best Local Similarity
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                                                                                                                                                 Query Match
Best Local Similarity
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                                                                                                                                   Matches
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TITLE OF INVENTION: Zinc Finger Domain Recognition
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CURRENT FILING DATE: 2003-01-23
PRIOR APPLICATION NUMBER: US 10/057,408
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                                                                                                                                                                                                                                                                                                                                                   LENGTH: 55
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OTHER INFORMATION: Nucleotides 28-30,
OTHER INFORMATION: 9, a, t, or c
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                                                              55 CTTTAGTNNNAGTNNNNNNTTACAANNNCACCAACGCACGCACACGGGGGAGAAG 1
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Pred. No. 2.4e+04;
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US-10-310-156-470/c
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Best Local
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                                                                                                                APPLICANT: Feldmann, Richard J.; Global Determinants, TITLE OF INVENTION: Chlamydophila pneumoniae AR39, correct Edward Control of the Contro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
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                                                              CURRENT APPLICATION NUMBER: US/10/310,156
CURRENT FILING DATE: 2002-12-05
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NUMBER OF SEQ ID NOS: 742 SOFTWARE: Proprietary
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TYPE: DNA
ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FEATURE:
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28; Conserv
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                                                                                                                                                    Richard J.; Global Determinants, Chlamydophila pneumoniae AR39, Co
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Pred. No. 2.
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Pred. No. 2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 8;
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; LOCATION: (40)...(40)
; OTHER INFORMATION: n is equal
US-09-912-293-732
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; LOCATION: (827553)...(827610)
; OTHER INFORMATION: Chromosome = 1
US-10-310-156-470
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                 Query Match
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NUMBER OF SEQ ID NOS:
SEQ ID NO 732
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Best Local Similarity
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LENGTH: 58
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PRIOR FILING DATE: 1993-08-09
PRIOR APPLICATION NUMBER: 08/196,363
PRIOR FILING DATE: 1994-02-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION NUMBER: US/09/912,293
CURRENT FILING DATE: 2001-07-26
PRIOR APPLICATION NUMBER: 08/103,744
PRIOR FILING DATE: 1993-08-09
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR FILING DATE: 1994-03-31 PRIOR APPLICATION NUMBER: 09/741,830
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                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR FILING DATE: 2001-03-21
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                                                                                                                               NAME/KEY: misc_feature
LOCATION: (38)..(38) OTHER INFORMATION: n is equal to
                                                                                                                                                                                                                    NAME/KEY: misc_feature LOCATION: (35)..(35)
                                                                                                                                                                                                                                                                                       NAME/KEY: misc_feature LOCATION: (29)..(29)
                                                                                                                                                                                                                                                                                                                      FEATURE:
                                                                                                                                                                                                                                                                                                                                        ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                       TYPE: DNA
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 Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: 09/813,155
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE: 2000-12-22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: 08/196,362
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE: 1994-03-31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: 08/221,623
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILING DATE: 1994-02-15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER: 09/859,490
2001-05-18
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Score 23; DB 6;
Pred. No. 4.2e+04;
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Pred. No. 2
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Sequence 48, Application PC/TUS0302358 GENERAL INFORMATION:

APPLICANT: Sera, Takashi

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RESULT 35
PCT-US03-02358-48/c
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Best Local Similarity
Watches 29; Conserva
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LENGTH: 60
TYPE: DNA
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Best Local S
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                                                                                                                                                                                                                                                                                                                                                                                     PRIOR FILING DATE: 2002-01-23
NUMBER OF SEQ ID NOS: 87
SOFTWARE: PatentIn version 3.2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Sera, Takashi
TITLE OF INVENTION: Zinc Finger Domain Recognition Code and Uses Thereof
FILE REFERENCE: 199845.158
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: PCT/US03/02358 CURRENT FILING DATE: 2003-01-23
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TYPE: DNA
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CURRENT FILING DATE: 2003-02-20
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TITLE OF INVENTION: NUCLEOTIDE AND AMINO ACID SEQUENCES
TITLE OF INVENTION: RELATING TO RESPIRATORY DISEASES AN
FILE REFERENCE: HUMO2-11P
                                                                                                                                                                                                              NAME/KEY: misc_feature
LOCATION: (37)..(51)
OTHER INFORMATION: Nucleotides 37-39 and
                                                                                                                                                                                                                                                                  FEATURE:
                                                                                                                                                                                                                                                                                  OTHER INFORMATION: Partial zinc finger domain oligomer
                                                                                                                                                                                                                                                                                                     ORGANISM: Artificial Sequence FEATURE:
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                                                                                 1239 CACCAGAGGGCGCACACTGGGGGAGAAGCCTTA 1270
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                                                                     36
                                                                                                                                l Similarity 81.2
26; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2 ACCCTTATCGAACATCAGCGAATTCACACTGGACAAAAAC
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                                                                  CATCAGCGTACTCACACTGGCGAGAAGCCTTA 5
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                                                                                                                                                0.8%;
                                                                                                                                                Score 22.4;
Pred. No. 6
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US-09-911-261B-48/c
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APPLICANT: Sera, Takashi
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SEQ ID NO 39
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   SEQ ID NO 48
                                                                TITLE OF INVENTION: Zinc Finger Domain Recognition Code and Uses Thereof FILE REFERENCE: 109845.135
CURRENT APPLICATION NUMBER: US/09/911,261B
CURRENT FILING DATE: 2000-07-21
PRIOR APPLICATION NUMBER: US 60/220,060
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PRIOR FILING DATE: 2002-01-23
NUMBER OF THE PRIOR FILING PATE: 2002-01-23
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TITLE OF INVENTION: Zinc Finger Domain Recognition Code and Uses Thereof
FILE REFERENCE: 109845.135
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CURRENT APPLICATION NUMBER: PCT/US03/02358
CURRENT FILING DATE: 2003-01-23
                                    PRIOR FILING DATE: 2000-
NUMBER OF SEQ ID NOS: 69
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NUMBER OF SEQ ID NOS: 87
                    SOFTWARE: PatentIn version 3.0
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les 26; Conserv
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; OTHER INFORMATION: Nucleotides
US-09-911-261A-39
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                                                     US-09-911-261A-48
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                                                                                                                                                        NUMBER OF SEQ ID NOS: 69
SOFTWARE: PatentIn version
SEQ ID NO 48
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LENGTH: 60
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CURRENT APPLICATION NUMBER: US/09/911,261A
CURRENT FILING DATE: 2001-07-23
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                                                                                                      ORGANISM: Artificial
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                                                                   OTHER INFORMATION: Partial zinc finger domain oligomer
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No. 6.3e+04;
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; ORGANISM: Mus musculus
US-60-427-808-134357
RESULT 42
US-60-427-836-480873
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; OTHER INFORMATION: Accession number cg42532030
US-09-749-280B-6410
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Xue Mei Zhou
TITLE OF INVENTION: Methods of Genetic Analysis of Mouse
FILE REFERENCE: 3528
CURRENT APPLICATION NUMBER: US/60/427,808
CURRENT FILING DATE: 2002-11-20
NUMBER OF SEQ ID NOS: 982914
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQ ID NOS: 10038
SOFTWARE: CuraGen Patent Fo:
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CURRENT FILING DATE: 2000-12-27
PRIOR APPLICATION NUMBER: 60/173,419
PRIOR FILING DATE: 1999-12-28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Shimkets, Richard A. APPLICANT: Leach, Martin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH: 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME/KEY: misc_feature
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FEATURE:
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64.7%;
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Pred. No. 6.3e+04;
0; Mismatches 2
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Pred. No. 6.7e+04;
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PCT-US03-11497-11
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GENERAL INFORMATION:
APPLICANT: Keith, Tim
TITLE OF INVENTION: NUCLEOTIDE AND AMINO ACID SEQUENCES
TITLE OF INVENTION: RELATING TO RESPIRATORY DISEASES AN
FILE REFERENCE: HUMO2-11P
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SEQ ID NO 517
LENGTH: 41
TYPE: DNA
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SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1
SEQ ID NO 480873
LENGTH: 25
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Best Local
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                                                                                                                                                                                  PRIOR APPLICATION NUMBER: US 60/372,669
PRIOR FILING DATE: 2002-04-16
PRIOR APPLICATION NUMBER: US 60/411,882
PRIOR FILING DATE: 2002-09-20
                                                                                                PRIOR APPLICATION NUMBER: US 60/424,336
PRIOR FILLING DATE: 2002-11-07
PRIOR APPLICATION NUMBER: US 60/374,823
PRIOR FILLING DATE: 2002-04-24
                                                                                                                                                                                                                                                                      CURRENT APPLICATION NUMBER: PCT/US03/11497 CURRENT FILING DATE: 2003-04-15
                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: TISSUE :
FILE REFERENCE: 16U 200 PCT
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TITLE OF INVENTION: TISSUE SPECIFIC GENES AND GENE CLUSTERS
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CURRENT FILING DATE: 2003-02-20
NUMBER OF SEQ ID NOS: 1000
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CURRENT FILING DATE: 2002-11-20
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TITLE OF INVENTION: Methods
FILE REFERENCE: 3527
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                                      FILING DATE: 2002-05-01 APPLICATION NUMBER: US 60/381,366
                                                                               APPLICATION NUMBER: US 60/376,558
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23; Conserv
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NUMBER: US 60/403,648
                       2002-05-
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Pred. No. 7.9e+04;
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CURRENT FILING DATE: 2002-10-17
PRIOR APPLICATION NUMBER: PCT/JP01/03399
PRIOR FILING DATE: 2001-04-20
PRIOR APPLICATION NUMBER: JP 12-126340
PRIOR APPLICATION NUMBER: JP 12-205587
PRIOR APPLICATION NUMBER: JP 12-205587
PRIOR APPLICATION NUMBER: JP 12-247962
PRIOR APPLICATION NUMBER: JP 12-247962
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                                  APPLICANT: Fry, Kirk
APPLICANT: Ly, Ngoc
APPLICANT: Woodward, Robert
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR DIAGNOSING AND MONITORING TRANSPLANT
TITLE OF INVENTION: REJECTION
FILE REFERENCE: 506612000122
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: OGI, Kazuhiro TITLE OF INVENTION: Novel Insulin/IGF/Relaxin Family Polypeptide and Its FILE REFERENCE: 2717 USOP
                                                                                                                                    APPLICANT: Wohlgemuth, Jay APPLICANT: Fry, Kirk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR FILING DATE: 2002-08-16
NUMBER OF SEQ ID NOS: 344
SOFTWARE: PatentIn version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR FILING DATE: 2000-08-10 PRIOR APPLICATION NUMBER: JP 12-395050 PRIOR FILING DATE: 2000-12-22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT:
CURRENT APPLICATION NUMBER: US/10/325,899
CURRENT FILING DATE: 2002-12-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FEATURE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 50
                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity
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                                                                                                                                                                                                                                                                                                                    577 ATCCAGAAGATCAGAAACAGCAGCAGGATCCATTCTGCTTTAGTGGCAA 625
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 32;
                                                                                                                                                                                                                                                                                              58 ACCCATACCGCCAGGAGCAGCAGCATGTATCTTGCCATGGTGGCAA 10
                                                                                                                                                                                                                                                                                                                                                                            32;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQ ID
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HARADA, Masataka
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NISHI, Kazunori
                                                                                                                                                                                                                                                                                                                                                                            Conservative
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Pred. No. 9.4e+04;
***cmatches 17;
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Pred. No. 8.6e+04;
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APPLICANT:
APPLICANT:

Woodward,

Robert

APPLICANT: Wohlgemuth, APPLICANT: Fry, Kirk APPLICANT: Ly, Ngoc

Jay

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RESULT 48
US-10-325-899-4844/c
; Sequence 4844, Application US/10325899
; GENERAL INFORMATION:
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; OTHER INFORMATION: Accession number cg44911411
US-09-749-280B-5462
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; Sequence 5462, Application US/09749280B
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US-10-325-899-4332
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SOFTWARE: CuraGen Patent Formatter Version 0.9
SEQ ID NO 5462
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Best Local
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                                                                                                                                                                                      Matches
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Shimkets, Richard A. APPLICANT: Leach, Martin
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NUMBER OF SEQ ID NOS: 9966
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: DNA
                                                                                                                                                                                                                                                                                                            NAME/KEY: misc_feature
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                                                                                                                                                                                                                                                                                                                                                   LOCATION: (25)...(26)
OTHER INFORMATION: Nucleotide
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LOCATION: (25)...(26)
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OTHER INFORMATION: single nucleotide polymorphism
                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME/KEY: allele
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27; Conserv
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75.0%;
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Pred. No. 1
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Pred. No. 9.9e+04;
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1.1e+05;
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RESULT 50
US-10-089-560-10/c
- Sequence 10, Application US/10089560
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: LOCATION: (0)...(0)

: OTHER INFORMATION: Accession number cg42924171

US-09-749-280B-1151
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Sequence 1151, Application US/09749280B
GENERAL INFORMATION:
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US-10-325-899-4844
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Best Local :
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TITLE OF INVENTION: Nucleic Acids Containing Single
FILE OF INVENTION: Nucleic Acids Containing Single
FILE REFERENCE: 15966-510
CURRENT APPLICATION NUMBER: US/09/749,280B
CURRENT FILING DATE: 2000-12-27
PRIOR APPLICATION NUMBER: 60/173,419
PRIOR FILING DATE: 1999-12-28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 50
TYPE: DNA
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NUMBER OF SEQ ID NOS: 10038
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION NUMBER: US 10/006,290 PRIOR FILING DATE: 2001-10-22 PRIOR APPLICATION NUMBER: US 10/131,831 PRIOR FILING DATE: 2002-04-24
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OTHER INFORMATION: Nucleotide deleted between bases
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PRIOR FILING DATE: 2001-06-08
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                                                                                                                                                                                                                                                                                                                                                    FEATURE
                                                                                                                                                                                                                                                                                                                                                                                                                                    OTHER INFORMATION: single nucleotide polymorphism
                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME/KEY: allele
LOCATION: (26)..
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26; Conserv
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71.8%;
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Pred. No. 1.1e
0; Mismatches
                                                                                                                                                                                                               Score 21.2; DB 5;
Pred. No. 1.3e+05;
                                                                                                                                                                                               Pred. No. 1.3e
); Mismatches
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1.1e+05;
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                                                                                                                                                                                                                                  Length 50;
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Query Match
Best Local Similarity
Thes 32; Conserve
Job time : 1950 secs
              Search completed: June 4,
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                                                                                                                                                                                                                                                                                                                     SEQ ID NO 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQ ID NOS: 10 , SOFTWARE: PatentIn version 3.1
                                                                                                                                                                                                                                                                                                                                                                       CURRENT FILING DATE: 2002-04-01
PRIOR APPLICATION NUMBER: 60/158,664
PRIOR FILING DATE: 1999-10-08
                                                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION NUMBER: US/10/089,560 CURRENT FILING DATE: 2002-04-01
                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Liu, Gang
TITLE OF INVENTION: Particle Analysis Assay for Biomolecular Quantification
FILE REFERENCE: 1321.2.71
                                                                                                                                                                                                                                                                                    LENGTH: 50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Straume, Tore APPLICANT: Liu, Gang
                                                                                                                                                                                                                                                          ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                 FEATURE:
                                                                                   1592 GGGGGAGAAGCCATTTGTATGTATGTACGGAGTGTGGGCGAGGCTTTACCCGGA 1641
                                                                50 GTGGAAGATGCCCTTTGAAGGCATGGGGTTTGGGTTAGATTTTTGTCTGA 1
                                                                                                                                         Conservative
                                                                                                                                                         0.8%;
              2003, 12:40:27
                                                                                                                                   Score 21.2; DI Pred. No. 1.3e-0; Mismatches
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1.3e+05;
hes 18;
                                                                                                                                                                      Length 50;
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Post-processing: Minimum Match 0%
Maximum Match 100%
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Maximum DB
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Title:
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2772
1 caggcgcgttaagctggttg.....tttctacccatcctcaccct 2772
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Gapop 10.0 , Gapext 1.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GenCore version 5.1.6 Copyright (c) 1993 - 2003 Compugen Ltd
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

Result No.

Score

Query Match Length

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                                                                 TITLE OF INVENTION: OLIGONUCLECTIDE LIBRARY FOR DETECTING RNA TRANSCRIPTS AND TITLE OF INVENTION: THAT POPULATE A TRANSCRIPTOME
FILE REFERENCE: 36688-0005
CURRENT APPLICATION NUMBER: US/09/908,975
CURRENT FILING DATE: 2001-07-20
PRIOR APPLICATION NUMBER: US 60/287,724
PRIOR PELICATION NUMBER: US 60/287,724
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PRIOR FILING DATE: 2001-05-02
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US-60-213-360-2913
US-60-213-360-2913
US-60-213-360-2913
US-60-213-360-2913
US-60-250-830-1005
US-60-250-830-1005
US-60-250-830-1005
US-60-278-232-10137
US-60-278-258-258-2006-1005
US-60-278-258-258-2006-1005
US-60-323-966-1005
US-60-323-966-1005
US-60-323-966-1106
US-60-323-966-1106
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US-60-323-966-1005
US-60-323-966-1005
US-60-323-966-1006
US-60-235-536-536-5360
US-60-235-536-5360
US-60-255-5374A-13100
US-60-172-373-18177
Score
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2913, Ap
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APPLICANT: FAIGLE, SIMCHON
TITLE OF INVENTION: OLIGONUCLEOTIDE LIBRARY FOR DETECTING RNA
TITLE OF INVENTION: THAT POPULATE A TRANSCRIPTOME
FILE REFERENCE: 36688-0004
CURRENT APPLICATION NUMBER: US/60/287,724
CURRENT FILING DATE: 2001-05-02
NUMBER OF SEQ ID NOS: 32337
SOFTWARE: Patentin version 3.0
SEQ ID NO 6782
LENGTH: 60
TYPE: DNA
ORGANISM: Homo sapiens
US-60-287-724-6782
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION NUMBER: US/09/908,975A
CURRENT ETILING DATE: 2001-07-20
PRIOR APPLICATION NUMBER: US 60/287,724
PRIOR FILING DATE: 2001-05-02
PRIOR APPLICATION NUMBER: US 60/221,607
PRIOR APPLICATION NUMBER: US 60/221,607
PRIOR FILING DATE: 2000-07-28
NUMBER OF SEQ ID NOS: 32337
SOFTWARE: PATENTIN VETSION 3.0
SEQ ID NO 6782
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US-09-908-975A-6782
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US-60-287-724-6782
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TYPE: DNA
ORGANISM: Homo sapiens
US-09-908-975A-6782
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; GENERAL INFORMATION:
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APPLICANT: MINTZ, Liat
APPLICANT: FAIGLER, Sinchon
APPLICANT: FAIGLER, Sinchon
TITLE OF INVENTION: OLIGONUCLEOTIDE LIBRARY FOR DETECTING RNA TRANSCRIPTS AND SPLI
TITLE OF INVENTION: THAT POPULATE A TRANSCRIPTOME
FILE REFERENCE: 36688-0006
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APPLICANT: WASSERMAN,
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APPLICANT: MINTZ, Liat
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Similarity 100.0%; Pred. No.
60; Conservative 0; Mismatc
                                                                                   Similarity
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MINTZ, Eli
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                                                                   Score 60; DB 72;
; Pred. No. 3.4e-06;
0; Mismatches 0;
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thes 0;
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: LOCATION: 26
: OTHER INFORMATION: 346325.1, 757, T->C
US-60-172-373-18032
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LOCATION: 26
: OTHER INFORMATION: 346325.1, 410, G->T
US-60-172-373-18031
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US-60-172-373-18031
                                                                                          Query Match
Best Local
                                                                         Matches
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SEQ ID NO 18032
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CURRENT FILING DATE: 1999-12-16
NUMBER OF SEQ ID NOS: 25,772
SOFTWARE: PERL Program
SEQ ID NO 18031
LENGTH: 51
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Best Local
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APPLICANT: Diep, Dinh
TITLE OF INVENTION: Method for the Identification of S
TITLE OF INVENTION: Polynucleotide Sequence Databases,
EILE REFERENCE: GX-0006 p
CURRENT FILLING DATE: 1999-12-16
CURRENT FILLING DATE: 1999-12-16
CURRENE SEEL DANS: 25,772
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GENERAL INFORMATION:
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APPLICANT: Diep, Dinh
TITLE OF INVENTION: Method for the Identification of Sequence Polymorphisms Using
TITLE OF INVENTION: Polynucleotide Sequence Databases, and Single Nucleotide Polyn
ETLE REFERENCE: GX-0006 P
                                                                                                                                                                                                                NAME/KEY: misc_feature
OTHER INFORMATION: Incyte ID No: SNP00003873
                                                                                                                                                                                                                                                                     ORGANISM: Homo sapiens
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                                                                                                                                                                                                                                                                                                           ENGTH: 51
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                                                                                        Local Similarity
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                             1974 TCACACCTCATTAGACACCAGAGGACACATTCAGGAGAGAGCCTTATATT 2024
                                                                                                                                                                                                                                                                                               DNA
Similarity
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100.0%;
                                                                  Score 51; DB; Pred. No. 0.0
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Pred. No.
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                                                                  DB 61; I
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; NAME/KEY: misc_feature
; LCCATION: (0)...(0)
; OTHER INFORMATION: Accession number cg43297259
US-09-718-321A-869
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                                                                                                                                                                                                                                                                    SEQ ID NO 869
LENGTH: 51
TYPE: DNA
                         Query Match
Best Local
     Matches
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Best Local Similarity
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TITLE OF INVENTION: NUCLEIC ACIDS CONTAINING
TITLE OF INVENTION: USE THEREOF
FILE REFERENCE: 15966-599
                                                                                                                                                                                                                                                                                                                        SOFTWARE: CuraGen Patent Formatter Version 0.9
                                                                                                                                                                                                                                                                                                                                           PRIOR FILING DATE: 1999-11-24 NUMBER OF SEQ ID NOS: 1468
                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION NUMBER: 60/163,783
                                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION NUMBER: US/09/718,321A CURRENT FILING DATE: 2002-06-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Shimkets, Richard A. APPLICANT: Leach, Martin D.
                                                                                                                                                             FEATURE:
NAME/KEY: allele
LOCATION: (26)...(0)
OTHER INFORMATION: single nucleotide polymorphism
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TITLE OF INVENTION: NUCLEIC ACIDS CONTAINING SINGLE NUCLEIC ACID POLYMORPHISMS
TITLE OF INVENTION: USE THEREOF
                                                                                                                                                                                                                                              ORGANISM: Homo sapiens
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                                                                                                                                                       FEATURE
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OTHER INFORMATION: single nucleotide polymorphism
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l Similarity 98.0%;
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                   Score 49.4; DB 2
Pred. No. 0.0036;
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Pred. No. 0.0036;
Mismatches
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                                    DB 28;
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                                  Length 51;
Indels
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; OTHER INFORMATION: US-60-172-373-18033
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US-60-172-373-18033
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LOCATION: 26
LOCATION: 07
JUSTINFORMATION: 346325.1,
US-60-278-258-5294
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-60-278-258-5294
                                                                                                                                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/60/278,258
CURRENT FILING DATE: 2001-03-23
NUMBER OF SEQ ID NOS: 17730
SOFTWARE: PERL Program
SEQ ID NO 5294
LENGTH: 51
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LENGTH: 51
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APPLICANT: Diep, Dinh
TITLE OF INVENTION: Method for the Identification of Sequence Polymorphisms Using
TITLE OF INVENTION: Polynucleotide Sequence Databases, and Single Nucleotide Polymor
FILE REFERENCE: GX-0006 P
                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Diep, Dinh
TITLE OF INVENTION: Method for the Identification of Sequence Polymorphisms Using
TITLE OF INVENTION: Polymucleotide Sequence Databases, and Single Nucleotide
TITLE OF INVENTION: Polymorphisms Identified Thereby
FILE REFERENCE: GX-0010-1 P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION NUMBER: US/60/172,373
CURRENT FILLING DATE: 1999-12-16
NUMBER OF SEQ ID NOS: 25,772
SOFTMARE: PERL Program
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                                                                                                                                                                                                    NAME/KEY: misc_feature
OTHER INFORMATION: Inc
                                                                                                                                                                                                                                     FEATURE:
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LOCATION: 26
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME/KEY: misc_feature
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: DNA
ORGANISM: Homo sapiens
                                                                                 Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        643 AGGGAGAAGACTCCAGACTCCTGTTTGGGAGAGTAAGCAAAAATGGCACTT 693
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity
50; Conser
                                                                                   Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AGGGAGTGTGAGCAAGGCTTTAGCCGGAAGTCACACCTCATCAGACACTTA 1
   AGGGAGAAGACTCCAGACTCCTGTTGGGGAGAGTAAGCAAAAATGGCACTT 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Lal, Preeti
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1.8%;
nilarity 98.0%;
Conservative
                                                                   Conservative
                                                                                                                                                                                                    Incyte ID
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                                                                                 1.8%;
                                                                                                                                                   421,
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                                                                                                                                                                                                    No: SNP00075163
                                                                                 Score 49.4; DB 7
Pred. No. 0.0036;
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Pred. No. 0
                                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
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US-60-278-258-5296
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Matches :
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                                                                                                 Query Match
Best Local
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FEATURE:
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LOCATION: 26
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                                                                                                                                                                                                                                                                                                  LENGTH: 51
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Sequence 5296, Appli GENERAL INFORMATION:
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APPLICANT: Diep, Dinh
TITLE OF INVENTION: Method for the Identification of Sequence Polymorphisms Using
TITLE OF INVENTION: Polymorphisms Identified Thereby
FILE REFERENCE: GX-0010-1 p
FILE REFERENCE: GX-0010-1 p
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Morris, MacDonald APPLICANT: Lal, Preeti APPLICANT: Diep, Dinh
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CURRENT FILING DATE: 2001-03-23
NUMBER OF SEQ ID NOS: 17730
                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/60/278,258
CURRENT FILING DATE: 2001-03-23
NUMBER OF SEQ ID NOS: 17730
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        THILE OF INVENTION: Method for the Identification of Sequence Polymorphisms Using THILE OF INVENTION: Polymorlectide Sequence Databases, and Single Nucleotide TITLE OF INVENTION: Polymorphisms Identified Thereby FILE REFERENCE: GX-0010-1 P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Morris, MacDonald
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME/KEY: misc_feature OTHER INFORMATION: Incyte ID No: SNP00075165
                                                                                                                                                                                    NAME/KEY: snp
LOCATION: 26
OTHER INFORMATION: 346325.1,
                                                                                                                                                                                                                                                       NAME/KEY: misc_feature OTHER INFORMATION: Incyte ID No: SNP00075166
                                                                                                                                                                                                                                                                                                                               ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     OTHER INFORMATION: 346325.1, 914, A->C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1136 GAAGTCGAACCTCTTTACACATCAGCGGACACACTCAGGGCTCAAGCCTTA 1186
2483 TTGGTGTGCTTTCCTCCGATTGATCCCAACCCTTCACCTATTTTACGTATA 2533
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 GAAGTCAAACCTCTTTACACATCAGCGGACACACTCAGGGCTCAAGCCTTA 51
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                                                                                           Similarity
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                                                                                           1.8%;
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                                                                                         Score 49.4; DB 71; Pred. No. 0.0036;
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US-50-278-258-5295
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                                                                     Best Local Similarity
Matches 49; Conserv
                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION NUMBER: US/60/278,258
CURRENT FILING DATE: 2001-03-23
NUMBER OF SEQ ID NOS: 17730
SOFTWARE: PERL Program
SEQ ID NO 5295
LENGTH: 51
                                                                                                          Query Match
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LENGTH: 51
TYPE: DNA
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APPLICANT: Diep, Dinh
TITLE OF INVENTION: Method for the Identification of Sequence Polymorphisms Using
TITLE OF INVENTION: Polymucleotide Sequence Databases, and Single Nucleotide
TITLE OF INVENTION: Polymorphisms Identified Thereby
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/60/324,185
CURRENT FILING DATE: 2001-09-21
NUMBER OF SEQ ID NOS: 35862
SOFTWARE: PERL Program
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                                                                                                                                                                                                            NAME/KEY: misc_feature OTHER INFORMATION: Incyte ID No: SNP00075164
                                                                                                                                                              OTHER INFORMATION: 346325.1,
                                                                                                                                                                                NAME/KEY: snp
LOCATION: 26
                                                                                                                                                                                                                                                      FEATURE:
                                                                                                                                                                                                                                                                 ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                         TYPE: DNA
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LOCATION: 26
OTHER INFORMATION: 346325.1,
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OTHER INFORMATION: Incyte ID No: SNP00115145
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity 98.0 ses 50; Conservative
                1108 GCAAGGATTGTGGACGAGGCTTTACTTGGAAGTCGAACCTCTTTACACATC 1158
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GCAAGGATTGTGGACGAGGCTTTACGTGGAAGTCAAACCTCTTTACACATC 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Application US/60278258
                                                                         Conservative
                                                                                      1.78; 96.18;
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                                                                                                                                                              , 988
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                                                                                    Score 47.8; DB Pred. No. 0.011;
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Pred. No. 0.0036;
                                                                     Mismatches
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                                                                                                      71;
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                                                                     Indels
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; LOCATION: 26
; OTHER INFORMATION: 1082325.11, 1102, C->T
US-60-324-185-4221
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                                                                                  Matches
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SEQ ID NO 4223
LENGTH: 51
TYPE: DNA
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CURRENT FILING DATE: 2001-09-21
NUMBER OF SEQ ID NOS: 35862
SOFTWARE: PERL Program
SEQ ID NO 4221
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Best Local Similarity
Matches 45; Conserv
                                                                                                       Best Local
                                                                                                                        Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Lal, Preeti
APPLICANT: Diep, Diah
TITLE OF INVENTION: METHOD FOR THE IDENTIFICATION OF SEQUENCE POLYMORPHISMS USING
TITLE OF INVENTION: POLYNUCLEOTIDE SEQUENCE DATABASES, AND SINGLE NUCLEOTIDE
TITLE OF INVENTION: POLYMORPHISMS IDENTIFIED THEREBY
FILE REFERENCE: GX-0019-1 P
                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION NUMBER: US/60/324,185
CURRENT FILING DATE: 2001-09-21
NUMBER OF SEQ ID NOS: 35862
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Diep, Dinh
TITLE OF INVENTION: METHOD FOR THE IDENTIFICATION OF SEQUENCE POLYMORPHISMS USING
TITLE OF INVENTION: POLYMORPHISMS EQUENCE DATABASES, AND SINGLE NUCLEOTIDE
TITLE OF INVENTION: POLYMORPHISMS IDENTIFIED THEREBY
FILE REFERENCE: GX-0019-1 P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Morris, MacDonald
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                                                                                                                                                                                                                                                     FEATURE:
NAME/KEY: misc_feature
                                                                                                                                                                                   OTHER INFORMATION: 1082325.7, 179, T->C
                                                                                                                                                                                                        NAME/KEY: snp
LOCATION: 26
                                                                                                                                                                                                                                              OTHER INFORMATION: Incyte ID No: SNP00097881
                                                                                                                                                                                                                                                                                                            ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME/KEY: misc_feature OTHER INFORMATION: Incyte ID No: SNP00106336
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ORGANISM: Homo sapiens
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                                      240 TTCACCCAGGAGGAGTGGAGGTTGTTGAGCCCTGCTCAGAGGACCCTGCA 289
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                                                                                  44;
2 TTCACCCAGGATGAGTGGAGGCTGCTGAGCCCTGCTCAAAGGACTCTGTA 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 GGACACACTCAGGGGAAAAGCCCTTATGTGTGCCGGGAATGTGGCAAAGGCT 51
                                                                                                   Similarity
                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Application US/60324185
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                                                                                                 Score 40.4;
Pred. No. 1
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                                                                                Mismatches
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RESULT 18
US-60-323-843-1769/c
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SEQ ID NO 20
LENGTH: 51
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SEQ ID NO 1769
LENGTH: 51
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                                                                                                                                           Query Match
Best Local Similarity
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Best Local
                                                                                                                               Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Lal, Preeti
APPLICANT: Diep, Dinh
TITLE OF INVENTION: POLYMORPHISMS USING TITLE OF INVENTION: POLYMORPHISMS USING TITLE OF INVENTION: POLYMORPHISMS IDENTIFIED THEREBY
TITLE OF INVENTION: POLYMORPHISMS IDENTIFIED THEREBY
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CURRENT FILING DATE: 2000-06-21
NUMBER OF SEQ ID NOS: 7924
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TITLE OF INVENTION: MG
TITLE OF INVENTION: PC
TITLE OF INVENTION: IC
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                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/60/234,690
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                                                                                                                                                                                                                                                      NAME/KEY: misc_feature
OTHER INFORMATION: Incyte
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LOCATION: 26
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                                                                                                                                                                                                                           NAME/KEY: snp
LOCATION: 26
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                                                                                                                                                                                                                                                                                                          ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                            TYPE: DNA
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                                                                                             2464 TGAGAACCTGTCTTCCCATTTGGTGTGCTTTCCTCCGATTGATCCCAACCC 2514
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               43;
                                                              51 TGAGAACTTGTCTTCCTGTTTGGCATGCTTTCCTCTGATTGGTCCCCACCC 1
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                                                                                                                               Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Method for the Identification of Sequence Polymorphisms Using Polynucleotide Sequence Databases, and Single Nucleotide Polymo
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Pred. No. 2.
                                                                                                                                           Score 38.2; D
Pred. No. 6.4;
                                                                                                                                                                                                          448, G->A
                                                                                                                             Mismatches
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Mismatches
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RESULT 20
US-60-213-359-2531
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LENGTH: 51
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Best Local
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APPLICANT: Diep, Dinh
TITLE OF INVENTION: METHOD FOR THE IDENTIFICATION OF SEQUENCE POLYMORPHISMS USING
TITLE OF INVENTION: POLYMORPHISMS EQUENCE DATABASES, AND SINGLE NUCLEOTIDE
TITLE OF INVENTION: POLYMORPHISMS IDENTIFIED THEREBY
FILE REFERENCE: GX-0021-1 P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILE REFERENCE: GX-0019-1 P
CURRENT APPLICATION NUMBER: US/60/324,185
CURRENT FILING DATE: 2001-09-21
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APPLICANT: Diep, Dinh
TITLE OF INVENTION: METHOD FOR THE IDENTIFICATION OF SEQUENCE POLYMORPHISMS USING
TITLE OF INVENTION: POLYMORPHISMS IDENTIFIED THEREBY
TITLE OF INVENTION: POLYMORPHISMS IDENTIFIED THEREBY
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CURRENT FILING DATE: 2001-09-21
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LOCATION: 26
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NAME/KEY: misc_feature
                                                                                                                                                                                                                                                                                                                                                    ORGANISM: Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                       TYPE: DNA
                                                                                                                                                                                                                                                           LOCATION: 26
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nes 41; Conserv
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                                                                         ACACACTCAGGGGAGAAACCTTTTGTTTGTAGAGAATGTGGGCAAGGATT 51
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Pred. No. 36;
0; Mismatches
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Sequence 2531, Application US/60213359

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                                                     RESULT 22
US-09-908-975A-6663
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: LOCATION: 26
: OTHER INFORMATION: 481631.7,
US-60-213-359-2531
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Sequence 6663, Application US/09908975A GENERAL INFORMATION:
APPLICANT: SHOSHAN, Avi
                                                                                                                                                                                                                Matches
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Best Local :
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LENGTH: 60
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SOFTWARE: PERL Progra
SEQ ID NO 2531
LENGTH: 51
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Best Local
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PRIOR FILING DATE: 2001-05-02
PRIOR APPLICATION NUMBER: US 60/221,607
PRIOR FILING DATE: 2000-07-28
NUMBER OF SEQ ID NOS: 32337
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CURRENT FILING DATE: 2001-07-20
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TITLE OF INVENTION: OLIGONUCLEOTIDE LIBRARY FOR DETECTING RNA TRANSCRIPTS AND SPLICE
TITLE OF INVENTION: THAT POPULATE A TRANSCRIPTOME
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CURRENT FILING DATE: 2000-06-21
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APPLICANT: Morris, MacDonald
                                                                                                                                                                                                                                                                                                                              TYPE: DNA
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OTHER INFORMATION: Incyte ID
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                                                                                                                                                                 1041 GGCTTTACGTGGAAGTCAAACCTGATCACACACAGAGGACACACTCAGGGGAGAAACC 1099
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MINTZ, Eli
MINTZ, Liat
                                                                                                                             GCCTTTTCCCAGAAGTCACACCTCACAGTACATCAGAGAATGCACACAGGGGAGAAACC
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                                                                                                                                                                                                      Score 35; DB 34; Length 60; Pred. No. 58; O; Mismatches 15; Indels
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PRIOR APPLICATION NUMBER: US 60/221,607
PRIOR FILING DATE: 2000-07-28
NUMBER OF SEQ ID NOS: 32337
SOFTWARE: Patentin version 3.0
SEQ ID NO 6663
LENGTH: 60
TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                              RESULT 24
US-09-726-173A-4584
Sequence 4584, Application US/09726173A
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
APPLICANT: Shimkets, Richard A.
APPLICANT: Leach, Martin D.
TITLE OF INVENTION: NUCLEIC ACIDS CONTAINING
TITLE OF INVENTION: USE THEREOF
FILE REFERENCE: 15966-600
CURRENT APPLICATION NUMBER: US/09/726,173A
CURRENT FILING DATE: 2002-03-08
PRIOR APPLICATION NUMBER: 60/168,138
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: FAIGLER, Simchon
TITLE OF INVENTION: OLIGONUCLEOTIDE LIBRARY FOR DETECTING RNA TRANSCRIPTS AND SPLI
TITLE OF INVENTION: THAT POPULATE A TRANSCRIPTOME
FILE REFERENCE: 36688-0004
CURRENT APPLICATION NUMBER: US/60/287,724
CURRENT APPLICATION NUMBER: US/60/287,724
CURRENT FILING DATE: 2001-05-02
NUMBER OF SEQ ID NOS: 32337
SOFTWARE: PRECEIT VERSION 3.0
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Best Local Similarity
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APPLICANT: WASSERMAN, Alon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT FILING DATE: 2001-07-20 PRIOR APPLICATION NUMBER: US 60/287,724 PRIOR FILING DATE: 2001-05-02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: FAIGLER, Simchon
TITLE OF INVENTION: OLIGONUCLEOTIDE LIBRARY FOR DETECTING RNA TRANSCRIPTS
TITLE OF INVENTION: THAT POPULATE A TRANSCRIPTOME
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILE REFERENCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT:
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                                                                                                                                                                                                                                                                                                      1041 GGCTTTACGTGGAAGTCAAACCTGATCACACACACAGAGGACACACTCAGGGGAGAAACC
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MINTZ, Liat
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; Pred. No. 58;
0; Mismatches
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58;
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RESULT 26
US-60-172-360-29152
; Sequence 29152 Application US/60172360
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US-60-324-185-4342
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Best Local S
Matches 39
                                                                                                                        GENERAL INFORMATION:
APPLICANT: Morris, MacDonald
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: PERL Program SEQ ID NO 4342
            APPLICANT: Lal, Preeti
APPLICANT: Diep, Dinh
TITLE OF INVENTION: Method for the Identification of So
TITLE OF INVENTION: Polynucleotide Sequence Databases,
FILE REFERENCE: GX-0007 P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION NUMBER: US/60/324,185
CURRENT FILING DATE: 2001-09-21
NUMBER OF SEQ ID NOS: 35862
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Lal, Preeti
APPLICANT: Diep, Dinh
APPLICANT: Diep, Dinh
TITLE OF INVENTION: METHOD FOR THE IDENTIFICATION OF SEQUENCE POLYMORPHISMS USING
TITLE OF INVENTION: POLYMUCLEOTIDE SEQUENCE DATABASES, AND SINGLE NUCLEOTIDE
TITLE OF INVENTION: POLYMORPHISMS IDENTIFIED THEREBY
FILE REFERENCE: GX-0019-1 P
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CURRENT APPLICATION NUMBER: US/60/172,360
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LOCATION: (0)...(0)
OTHER INFORMATION: Accession number cg43960167
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OTHER INFORMATION: 2 of 2 allelic variants (4583 is other entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: DNA
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ORGANISM: Homo sapiens
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LOCATION: 26
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                                                                                                                                                                                                                                                                                                                                                        Local Similarity es 39; Conserv
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                                                                                                                                                                                                                                                                                                           1056 TCAAACCTGATCACACATCAGAGGACACACTCAGGGGAGAAACCTTAT 1103
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Pred. No. 70;
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                                    Sequence Polymorphisms Using
s, and Single Nucleotide Polymor
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; NAME/KEY: misc_feature
, OTHER IMFORMATION: Incyte ID I
: NAME/KEY: snp
: LOCATION: 26
; OTHER INFORMATION: 1079415.2,
US-60-250-830-1043
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                                                                                                                                                                                                     RESULT 28
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Best Local Similarity
Matches 37; Conserv
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NUMBER OF SEQ ID NOS: 29838
SOFTWARE: PERL PROGRAM
SEQ ID NO 29152
                                                                                                                                               GENERAL INFORMATION:
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Best Local Similarity
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     APPLICANT: Lal, Preeti
APPLICANT: Diep, Dinh
APPLICANT: Diep, Dinh
TITLE OF INVENTION: METHOD FOR THE IDENTIFICATION OF SEQUENCE POLYMORPHISMS USING
TITLE OF INVENTION: POLYMORPHISMS IDENTIFIED THEREBY
FILE REFERENCE: GX-0020-1 P
FILE REFERENCE: GX-0020-1 P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Diep, Dinh
TITLE OF INVENTION: METHOD FOR THE IDENTIFICATION OF SI
TITLE OF INVENTION: POLYMUCLEOTIDE SEQUENCE DATABASES,
TITLE OF INVENTION: POLYMORPHISMS IDENTIFIED THEREBY
FILE REFERENCE: GX-0020 P
CURRENT APPLICATION NUMBER: US/60/250,830
CURRENT FILING DATE: 2000-11-04
NUMBER OF SEQ ID NOS: 3246
                                                                                                                            APPLICANT: Morris, MacDonald
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CURRENT APPLICATION NUMBER: US/60/323,966
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LOCATION: 26
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                                                                                                                                                                                                                                                                           229 TGGCTGTGTACTTCACCCAGGAGGAGGAGGTTGTTGAGCCCTGCTCAGA 279
                                                                                                                                                                                                                                                                                                                                 40;
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                                                                                                                                                                                                                                                          TGGCTGTGGACTTCACCCAGGAGGAATGGACTTTGCTGGATCAAACTCAGA 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PERL Program
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86.0%;
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Pred. No. 1.6e+02;
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Pred. No. 1.6e+02;
                                                                                                                                                                                                                                                                                                                                 Mismatches
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RESULT 30
US-60-172-373-5373
Sequence 5373, Application US/60172373
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                                                                                                                           GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                               Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                Matches
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SOFTWARE: CuraGen Patent Formatter Version 0.9
SEQ ID NO 4583
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 4583, Application US/09726173A GENERAL INFORMATION:
                APPLICANT: Morris, MacDonald
APPLICANT: Lal, Preeti
APPLICANT: Liep, Dinh
TITLE OF INVENTION: Method for the Identification of Sequence Polymorphisms Using
TITLE OF INVENTION: Polynucleotide Sequence Databases, and Single Nucleotide Polymorphisms FILE REFERENCE: GX-0006 P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT FILING DATE: 2001
NUMBER OF SEQ ID NOS: 3246
SOFTWARE: PERL Program
SEQ ID NO 1043
LENGTH: 51
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Best Local :
CURRENT APPLICATION NUMBER: US/60/172,373
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/09/726,173A CURRENT FILING DATE: 2002-03-08 PRIOR APPLICATION NUMBER: 60/168,138 PRIOR FILING DATE: 1999-11-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Leach, Martin D.
TITLE OF INVENTION: NUCLEIC ACIDS CONTAINING SINGLE NUCLEIC ACID POLYMORPHISMS AND
TITLE OF INVENTION: USE THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Shimkets, Richard A. APPLICANT: Leach, Martin D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILE REFERENCE: 15966-600
                                                                                                                                                                                                                                                                                                                                                                                               OTHER INFORMATION: Accession number cg43960167
                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME/KEY: misc_feature LOCATION: (26)...(0)
OTHER_INFORMATION: 1 of 2 allelic variants (4584)
                                                                                                                                                                                                                                                                                                                                                                                                                     NAME/KEY: misc_feature LOCATION: (0)...(0)
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NAME/KEY: snp
LOCATION: 26
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ORGANISM: Homo sapiens
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                         FEATURE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: DNA
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                                                                                                                                                                                                                                                    2375 AGTCAAATGCCCAGGCAGATAGGGGTGGGTACCTGGTGAAACCCAACCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             229 TGGCTGTGTACTTCACCCAGGAGGAGGTGGAGGTTGTTGAGCCCTGCTCAGA 279
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                                                                                                                                                                                                                                                                                                              Similarity 39; Conserv
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40; Conserv
                                                                                                                                                                                                                               AGACAAATGCCTAGGCAGATAGGGGCAGGTCAACAGTGAAACCCCCACCT 49
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79.6%;
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                                                                                                                                                                                                                                                                                                                           Score 33; DB 29
Pred. No. 2e+02;
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Pred. No. 1.6e+02;
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RESULT 32
US-60-278-258-5877
Sequence 5877, Application US/60278258
; GENERAL INFORMATION:
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: COCATION: 26
: OTHER INFORMATION: 414789.1,
US-60-172-373-5373
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: PER SEQ ID NO 2773 LENGTH: 51 TYPE: DNA
                                                                                                                                                                                                                                                                                                Best Local Similarity Matches 38; Conserv
                APPLICANT: Morris, MacDonald
APPLICANT: Lal, Preeti
APPLICANT: Lal, Preeti
APPLICANT: Diep, Dinh
TITLE OF INVENTION: Method for the Identification of Sequence Polymorphisms Using
TITLE OF INVENTION: Polymorphisms Identified Thereby
FILE REFERENCE: GX-0010-1 p
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Best Local Similarity
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NUMBER OF SEQ ID NOS: 25,772
SOFTWARE: PERL PROGRAM
SEQ ID NO 5373
CURRENT APPLICATION NUMBER: US/60/278,258
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CURRENT FILING DATE: 2000-06-21
NUMBER OF SEQ ID NOS: 8347
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TITLE OF INVENTION: Method for the Identification of Sequence Polymorphisms Using
TITLE OF INVENTION: Polynucleotide Sequence Databases, and Single Nucleotide Poly
TITLE OF INVENTION: Identified Thereby
FILE REFERENCE: GX-0014 P
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LOCATION: 26
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                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME/KEY: misc_feature
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: Homo sapiens
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                                                                                                                                                                                                                                                                                             Score 32.6; DB 65;
Pred. No. 2.7e+02;
D; Mismatches 9;
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RESULT 33
US-60-324-185-4224
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                                                                                                                                                        Sequence 6741, Application US/60324185 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION NUMBER: US/60/324,185
CURRENT FILING DATE: 2001-09-21
NUMBER OF SEQ ID NOS: 35862
SOFTWARE: PERL PROGRAM
SEQ ID NO 4224
LENGTH: 51
TYPE: DNA
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Best Local
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LENGTH: 51
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Best Local
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APPLICANT: MOTTIS, MacDonald
APPLICANT: Lal, Preeti
APPLICANT: Diep, Dinh
TITLE OF INVENTION: METHOD FOR THE IDENTIFICATION OF SEQUENCE POLYMORPHISMS USING TITLE OF INVENTION: POLYNUCLEOTIDE SEQUENCE DATABASES, AND SINGLE NUCLEOTIDE TITLE OF INVENTION: POLYNORPHISMS IDENTIFIED THEREBY
FILE REFERENCE: GX-0019-1 P
CURRENT APPLICATION NUMBER: US/60/324,185
CURRENT FILING DATE: 2001-09-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Lal, Preeti
APPLICANT: Diep, Dinh
TITLE OF INVENTION: METHOD FOR THE IDENTIFICATION OF SEQUENCE POLYMORPHISMS USING
TITLE OF INVENTION: POLYMORPHISMS EQUENCE DATABASES, AND SINGLE NUCLEOTIDE
TITLE OF INVENTION: POLYMORPHISMS IDENTIFIED THEREBY
TITLE REFERENCE: GX-0019-1 P
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NUMBER OF SEQ ID NOS: 17730
SOFTWARE: PERL Program
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OTHER INFORMATION: Incyte ID
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LOCATION: 26
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OTHER INFORMATION: Incyte ID No: SNP00065280
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LOCATION: 26
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                                                                                                                                                                                                                                                                                                                                               Local Similarity
                                                                                                                                                                                                                                                         1071 CATCAGAGGACACACTCAGGGGAGAAACCTTATGTGTGCAAGGATTGTGG 1120
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78.0%;
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                                                                                                                                                                                                                                                                                                                                           Pred.
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                                                                                                                                                                                                                                                                                                                                           4; DB 76;
3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            11;
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                                                                                                                                                                                                        RESULT 36
US-60-213-360-2926
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LOCATION: 26
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LOCATION: 26
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FEATURE:
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; OTHER INFORMATION: 332474.3, 556, G->A US-60-213-359-1622
                                                                                                                                                                    GENERAL INFORMATION:
APPLICANT: MOTTIS;
APPLICANT: Lal, Pr
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SOFTWARE: PERL Program
SEQ ID NO 6741
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                                                                                                                                                                                                                                        Sequence 2926, Application US/60213360
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APPLICANT: Diep, Dinh
TITLE OF INVENTION: Method for the Ident
TITLE OF INVENTION: Polynucleotide Seque:
TITLE OF INVENTION: Identified Thereby
FILE REFERENCE: GX-00114 P
CURRENT APPLICATION NUMBER: US/60/213,360
CURRENT FILING DATE: 2000-65-21
NUMBER OF SEQ ID NOS: 8347
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: Method for the Identification of Sequence Polymorphisms Using TITLE OF INVENTION: Polynucleotide Sequence Databases, and Single Nucleotide Poly TITLE OF INVENTION: Identified Thereby FILE REFERENCE: GX-0015 P CURRENT APPLICATION NUMBER: US/60/213,359 CURRENT APPLICATION NUMBER: US/60/213,359 CURRENT FILLING DATE: 2000-06-21 NUMBER OF SEQ ID NOS: 7924
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME/KEY: misc_feature
OTHER INFORMATION: Incyte ID No:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      OTHER INFORMATION: 1142546.1, 616, G->A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OTHER INFORMATION: Incyte ID No: SNP00118792
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                                                                                                                                                                                                                                                                                                                                                                                      276 CAGAGGACCCTGCACAGGGAGGTGATGCTGGAGACTTATAACCATCTG 323
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1622, Application US/60213359
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Lal, Preeti
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Morris, MacDonald
                                                                                                                                                                          Lal, Preeti
                                                                                                                                                                                                Morris; MacDonald
                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                            Method for the Identification Polynucleotide Sequence Databa
                                                                                                                                                                                                                                                                                                                                                                                                                                                       1.2%;
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78.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                   0:
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 32;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 32.4; DB 76; Pred. No. 3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         4e+02;
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                                                                                                            sequence Polymorphisms (
s, and Single Nucleotide
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RESULT 38
US-60-323-966-822
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                                                                                                                                                                                       Sequence 822, Application US/60323966 GENERAL INFORMATION:
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SEQ ID NO 822
LENGTH: 51
                                                                                                                                                                                                                                                                                                                                                                     Matches
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Best Local :
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                              APPLICANT: Diep, Dinh
TITLE OF INVENTION: METHOD FOR THE IDENTIFICATION OF SEQUENCE POLYMORPHISMS USING
TITLE OF INVENTION: POLYMORPHISMS EQUENCE DATABASES, AND SINGLE NUCLEOTIDE
TITLE OF INVENTION: POLYMORPHISMS IDENTIFIED THEREBY
FILE REFERENCE: GX-0020-1 P
CURRENT PEDLICATION NUMBER: US/60/323,966
CURRENT FILLING DATE: 2001-09-21
                                                                                                                                      APPLICANT: Morris, MacDonald APPLICANT: Lal, Preeti APPLICANT: Diep, Dinh
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NUMBER OF SEQ ID NOS: 3246 SOFTWARE: PERL Program
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/60/250,830
CURRENT FILING DATE: 2000-11-04
NUMBER OF SEQ ID NOS: 3246
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: METHOD FOR THE IDENTIFICATION OF SEQUENCE POLYMORPHISMS USING TITLE OF INVENTION: POLYMOCRETIDE SEQUENCE DATABASES, AND SINGLE NUCLECTIDE TITLE OF INVENTION: POLYMORPHISMS IDENTIFIED THEREBY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Lal, Preeti
APPLICANT: Diep, Dinh
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OTHER INFORMATION: Incyte ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME/KEY: snp
LOCATION: 26
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ORGANISM: Homo sapiens
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LOCATION: 26
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                                                                                                                                                                                                                                                                                                                                                                              Match 1.28;
Local Similarity 79.28;
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                                                                                                                                                                                                                                                                                    38;
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38; Conserv
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79.2%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       No: SNP00104016
                                                                                                                                                                                                                                                                                                                                                           Score 32; DB 69
Pred. No. 4e+02;
0; Mismatches
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Pred. No. 4e+02;
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                                                                                                                                                                                                                                                                                                                                                                                             DB 69; Length 51;
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RESULT 40
US-60-213-360-2750/c
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; LOCATION: 26
; OTHER INFORMATION: 482033.2, 1229, C->A
US-60-172-373-5588
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SOFTWARE:
                            APPLICANT: Morris, Macuouse APPLICANT: Lal, Preeti
APPLICANT: Lal, Preeti
APPLICANT: Diep, Dinh
TITLE OF INVENTION: Method for the Identification of Sequence Polymorphisms Using
TITLE OF INVENTION: Polynucleotide Sequence Databases, and Single Nucleotide Poly
TITLE OF INVENTION: Identified Thereby
TITLE OF INVENTION: Identified Thereby
                                                                                                                                                                                                                  GENERAL INFORMATION:
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APPLICANT: Diep, Dinh
TITLE OF INVENTION: Method for the Identification of Sequence Polymorphisms Using
TITLE OF INVENTION: Polynucleotide Sequence Databases, and Single Nucleotide Polym
FILE REFERENCE: GX-0006 p
FILE REFERENCE: GX-0006 p
CURRENT APPLICATION NUMBER: US/60/172,373
CURRENT APPLICATION NUMBER: US/60/172,373
CURRENT FILING DATE: 1999-12-16
NUMBER OF SEQ ID NOS: 25,772
SECTIMARE: PERL Program
SEQ ID NO 5588
                                                                                                                                                                                                                                   Sequence 2750, Application US/60213360
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Best Local (
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Best Local
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LENGTH: 51
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NAME/KEY: misc_feature
OTHER INFORMATION: Incy
FEATURE:
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TYPE: DNA
ORGANISM: Homo sapiens
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LOCATION: 26
OTHER INFORMATION: 332474.3,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME/KEY: misc_feature OTHER INFORMATION: Incyte ID No: SNP00104016
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                                                                                                                                                                                                                                                                                                                                           1200 TGTGGGCAGAGCTTTAGCCTGAAGTCAAACCTCATTACCCACCAGAGGGCG 1250
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                                                                                                                                                                                                                                                                                                                      1 TGTGGCCTGGGCTTCAGCGACAGGTAAAACCTCATCTCCCACCAGAGGACG 51
                                                                                                                                                                                                                                                                                                                                                                                                 39; Conservative
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38; Conserv
                 PERL Program
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79.2%;
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Pred. No. 4e+02;
0; Mismatches
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US-60-324-185-4044
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US-60-324-185-3926
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NUMBER OF SEQ ID NOS: 35862
SOFTWARE: PERL Program
SEQ ID NO 4044
LENGTH: 51
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GENERAL INFORMATION:
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Best Local
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                                                                                                                   TITLE OF INVENTION: POLYMUCLEOTIDE SEQUENCE DATABASE TITLE OF INVENTION: POLYMORPHISMS IDENTIFIED THEREBY FILE REFERENCE: GX-0019-1 P
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CURRENT FILING DATE: 2001-09-21
NUMBER OF SEQ ID NOS: 35862
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Diep, Dinh
TITLE OF INVENTION: METHOD FOR THE IDENTIFICATION OF SEQUENCE POLYMORPHISMS USING
TITLE OF INVENTION: POLYMUCLEOTIDE SEQUENCE DATABASES, AND SINGLE NUCLEOTIDE
TITLE OF INVENTION: POLYMORPHISMS IDENTIFIED THEREBY
                                                                                CURRENT APPLICATION NUMBER: US/60/324,185
CURRENT FILING DATE: 2001-09-21
                                                                                                                                     APPLICANT: Diep, Dinh
TITLE OF INVENTION: METHOD FOR THE IDENTIFICATION OF SEQUENCE POLYMORPHISMS USING
TITLE OF INVENTION: POLYNUCLEOTIDE SEQUENCE DATABASES, AND SINGLE NUCLEOTIDE
                                                                                                                                                                                                 APPLICANT:
APPLICANT:
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                                                                                                                                                                                                                                    APPLICANT: Morris, MacDonald
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OTHER INFORMATION: Incyte ID No: SNP00112420
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME/KEY: snp
LOCATION: 26
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OTHER INFORMATION: Incyte ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME/KEY: snp
LOCATION: 26
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: DNA
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Pred. No. 4.5e
0; Mismatches
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Pred. No. 5.
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ies 4;
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LOCATION: SOF
COTHER INFORMATION: 1079501.1,
US-60-324-185-4044
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; ORGANISM: Homo sapiens
US-09-908-975-6403
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US-09-908-975-6403
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Best Local Similarity
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CURRENT APPLICATION NUMBER: US/09/908,975
CURRENT FILING DATE: 2001-07-20
PRIOR APPLICATION NUMBER: US 60/287,724
PRIOR PILING DATE: 2001-05-02
PRIOR PILING DATE: 2000-07-28
PRIOR FILING DATE: 2000-07-28
                                                                                                                                APPLICANT: WASSERMAN, Alon
APPLICANT: MINTZ, Eli
APPLICANT: MINTZ, Liat
APPLICANT: FAIGLER, Simchon
TITLE OF INVENTION: OLIGONUCCLEOTIDE LIBRARY FOR DETECTING RNA TRANSCRIPTS AND
TITLE OF INVENTION: THAT POPULATE A TRANSCRIPTOME
FILE REFERENCE: 36688-0006
              PRIOR APPLICATION NUMBER: US 60/287,724
PRIOR TILING DATE: 2001-05-02
PRIOR APPLICATION NUMBER: US 60/221,607
PRIOR FILING DATE: 2000-07-28
                                                                                             CURRENT APPLICATION NUMBER: US/09/908,975A CURRENT FILING DATE: 2001-07-20
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TITLE OF INVENTION: OLIGONUCLEOTIDE LIBRARY FOR DETECTING RNA TRANSCRIPTS AND SPLI
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NUMBER OF SEQ ID NOS: 32337
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87.2%;
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77.68;
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Pred. No.
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Pred. No. 5.
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8.4e+02;
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ches 11;
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US-60-287-724-6403
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                                                                                                                                   CURRENT APPLICATION NUMBER: US/60/324,185
CURRENT FILING DATE: 2001-09-21
NUMBER OF SEQ ID NOS: 35862
SOFTWARE: PERL Program
SEQ ID NO 731
LENGTH: 51
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LENGTH: 60
TYPE: DNA
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APPLICANT: MORTIS, MacDonald
APPLICANT: Lal, Preeti
APPLICANT: Diep, Dinh
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APPLICANT: FAIGLER, Simchon
APPLICANT: OF INVENTION: OLIGONUCLEOTIDE LIBRARY FOR DETECTING RNA TRANSCRIPTS AND SPLICE
TITLE OF INVENTION: THAT POPULATE A TRANSCRIPTOME
FILE REFERENCE: 36688-0004
CURRENT APPLICATION NUMBER: US/60/287,724
CURRENT PILING DATE: 2001-05-02
NUMBER OF SEQ ID NOS: 32337
SOFTWARE: Patentin version 3.0
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                                                                                                                                                                                                                                      APPLICANT: Lal, Preeti
APPLICANT: Diep, Dinh
TITLE OF INVENTION: METHOD FOR THE IDENTIFICATION OF SEQUENCE POLYMORPHISMS USING
TITLE OF INVENTION: POLYNUCLEOTIDE SEQUENCE DATABASES, AND SINGLE NUCLEOTIDE
TITLE OF INVENTION: POLYNORPHISMS IDENTIFIED THEREBY
FILE REFERENCE: GX-0019-1 P
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OTHER INFORMATION: 017463.1, 155, T->C
                 NAME/KEY: snp
LOCATION: 26
                                               NAME/KEY: misc_feature
OTHER INFORMATION: Incyte
                                                                                     FEATURE
                                                                                                   ORGANISM: Homo sapiens
                                                                                                                      TYPE: DNA
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l Similarity 87.2%;
34; Conservative
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MINTZ, Liat
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87.2%;
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Pred. No. 8
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Pred. No. 8.4e+02;
                                               SNP00120539
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; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-953-198-448
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US-60-213-360-740
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APPLICANT: COMPUGEN LTD
TITLE OF INVENTION: DATA BASE OF SNP
FILE REFERENCE: 1351253
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APPLICANT:
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Best Local Similarity
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CURRENT FILING DATE: 2000-06-21
NUMBER OF SEQ ID NOS: 8347
                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION: Method for the Identification of Sequence Polymorphisms Using TITLE OF INVENTION: Polynucleotide Sequence Databases, and Single Nucleotide PolyTITLE OF INVENTION: Identified Thereby
                                                                                                                                                                                                                                                                                                                                                                          FILE REFERENCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/09/953,198
CURRENT FILING DATE: 2001-08-12
NUMBER OF SEQ ID NOS: 979
                                                                                                                                            OTHER INFORMATION: 1077201.1, 925, T->C
                                                                                                                                                                                            NAME/KEY: misc_feature
OTHER INFORMATION: Incyte ID No: SNP00078224
                                                                                                                                                                                                                                                             LENGTH: 51
TYPE: DNA
                                                                                                                                                             NAME/KEY: snp
LOCATION: 26
                                                                                                                                                                                                                             FEATURE:
                                                                                                                                                                                                                                            ORGANISM:
                1675 CAGGGGAGAAGCCATTTGTATGTGCTGAGTGTGGACGAGGCTTTA 1719
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               274 CTCAGAGGACCCTGCACAGGGAGGTGATGCTGGAGACTT
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                                                                               Similarity
CTGGGGAGAAGCCTTTTGAATGTGCGGAGTGTGGAAAATCCTTCA 46
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Lal, Preeti
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                                                             Score 30.6; DB 65; Pred. No. 1e+03; 0; Mismatches 9;
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Pred. No. 8.8e+02;
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NAME/KEY: misc_feature
OTHER INFORMATION: Incyte ID No: SNP00120410
: NAME/KEY: SNP
LOCATION: 26
: OTHER INFORMATION: 1080820.1, 293, T->G
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Search completed: June 4, Job time: 6069 secs
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LICATION: 26
COTHER INFORMATION: 1080820.1, 293, T->G
US-60-323-966-971
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CURRENT FILING DATE: 2001-09-21
NUMBER OF SEQ ID NOS: 3246
SOFTWARE: PERL Program
SEQ ID NO 971
LENGTH: 51
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CURRENT APPLICATION NUMBER: US/60/250,830
CURRENT FILING DATE: 2000-11-04
NUMBER OF SEQ ID NOS: 3246
SOFTWARE: PERL PROGRAM
SEQ ID NO 971
LENGTH: 51
                                                                                                                                                               Query Match 1.1%;
Best Local Similarity 80.0%;
Matches 36; Conservative
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APPLICANT: Diep, Dinh
APPLICANT: Diep, Dinh
TITLE OF INVENTION: METHOD FOR THE IDENTIFICATION OF SEQUENCE POLYMORPHISMS USING
TITLE OF INVENTION: POLYMUCLEOTIDE SEQUENCE DATABASES, AND SINGLE NUCLEOTIDE
TITLE OF INVENTION: POLYMORPHISMS IDENTIFIED THEREBY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: MORTIS, MacDonald
APPLICANT: Lal, Preeti
APPLICANT: Lal, Preeti
APPLICANT: Lide, Dinh
TITLE OF INVENTION: METHOD FOR THE IDENTIFICATION OF SEQUENCE POLYMORPHISMS USING
TITLE OF INVENTION: POLYMOCLECTIDE SEQUENCE DATABASES, AND SINGLE NUCLECTIDE
TITLE OF INVENTION: POLYMORPHISMS IDENTIFIED THEREBY
FILE REFERENCE: GX-0020-1 P
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                                                                                                                                                                                                                                                                                                                       NAME/KEY: misc_feature
OTHER INFORMATION: Incyte ID No: SNP00120410
                                                                                                                                                                                                                                                                                                                                                                    TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: DNA ORGANISM: Homo sapiens FEATURE:
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Local Similarity 80.0%;
hes 36; Conservative
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                   2003; 12:07:57
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Pred. No. 1e+03;
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| 227 18.2 0.7 46 9 US-09-997-542-239 228 18.2 0.7 46 9 US-09-990-427-239 229 18.2 0.7 46 9 US-09-990-439-239 230 18.2 0.7 46 9 US-09-989-328-239 231 18.2 0.7 46 9 US-09-981-323-239 232 18.2 0.7 46 9 US-09-941-992-239 233 18.2 0.7 46 9 US-09-941-992-239 234 18.2 0.7 46 9 US-09-992-521-239 235 18.2 0.7 46 9 US-09-997-333-239 236 18.2 0.7 46 9 US-09-997-333-239 237 18.2 0.7 46 10 US-09-989-722-239 238 18.2 0.7 46 10 US-09-989-722-239 239 18.2 0.7 46 10 US-09-989-723-239 230 18.2 0.7 46 10 US-09-989-723-239 231 18.2 0.7 46 10 US-09-989-723-239   | 210 18.2 0.7 46 9 US-09-997-601-239 218 18.2 0.7 46 9 US-09-997-601-239 219 18.2 0.7 46 9 US-09-998-729A-239 220 18.2 0.7 46 9 US-09-998-844-239 221 18.2 0.7 46 9 US-09-997-840-239 221 18.2 0.7 46 9 US-09-997-349-239 221 18.2 0.7 46 9 US-09-997-628-239 223 18.2 0.7 46 9 US-09-997-628-239 224 18.2 0.7 46 9 US-09-997-628-239 225 18.2 0.7 46 9 US-09-997-683-239 226 18.2 0.7 46 9 US-09-993-689-239 227 18.2 0.7 46 9 US-09-993-689-239 228 18.2 0.7 46 9 US-09-993-689-239   | 206 18.2 0.7 46 9 US-09-990-711-239 207 18.2 0.7 46 9 US-09-998-726-239 208 18.2 0.7 46 9 US-09-998-126-239 209 18.2 0.7 46 9 US-09-998-126-239 210 18.2 0.7 46 9 US-09-991-157-239 211 18.2 0.7 46 9 US-09-991-172-239 212 18.2 0.7 46 9 US-09-991-172-239 213 18.2 0.7 46 9 US-09-997-514-239 214 18.2 0.7 46 9 US-09-997-513-239 215 18.2 0.7 46 9 US-09-997-523-239 216 18.2 0.7 46 9 US-09-997-523-239 217 18.2 0.7 46 9 US-09-997-523-239  | 186     18.2     0.7     25     10     US-09-866-108-14629       187     18.2     0.7     34     10     US-09-730-857-30       188     18.2     0.7     43     10     US-09-744-420-2       189     18.2     0.7     43     12     US-10-082-032-2       190     18.2     0.7     45     10     US-09-760-008A-6       191     18.2     0.7     46     9     US-09-989-755-239       192     18.2     0.7     46     9     US-09-989-735-239       193     18.2     0.7     46     9     US-09-989-735-239       194     18.2     0.7     46     9     US-09-989-735-239       195     18.2     0.7     46     9     US-09-989-730-239       196     18.2     0.7     46     9     US-09-989-730-239       197     18.2     0.7     46     9     US-09-990-436-239       198     18.2     0.7     46     9     US-09-991-181-239       199     18.2     0.7     46     9     US-09-991-867-239       198     18.2     0.7     46     9     US-09-991-867-239       199     18.2     0.7     46     9     US-09-997-  | 171 18.6 0.7 46 10 US-09-738-847-21 172 18.6 0.7 50 9 US-09-754-066.15 173 18.6 0.7 51 10 US-09-826-630-7 18.6 0.7 51 10 US-09-826-630-8 175 18.6 0.7 51 10 US-09-826-630-8 175 18.6 0.7 55 9 US-09-991-209-72 176 18.6 0.7 55 9 US-09-991-209-72 176 18.4 0.7 56 7 US-08-781-986A-5140 177 18.4 0.7 38 9 US-09-931-325A-186 178 18.4 0.7 38 9 US-09-730-2898-2508 179 18.4 0.7 45 9 US-09-8730-2898-2508 179 18.4 0.7 48 9 US-09-8781-986A-616 181 18.4 0.7 59 10 US-09-878-944-9 182 18.4 0.7 59 10 US-09-8781-984-1803 183 18.4 0.7 60 9 US-09-979-999-19 185 18.4 0.7 60 9 US-09-979-999-19 186 18.4 0.7 60 9 US-09-979-999-19   | 166 18.8 0.7 49<br>167 18.8 0.7 51<br>168 18.6 0.7 25<br>169 18.6 0.7 38<br>170 18.6 0.7 42   |
| 227 18.2 0.7 46 9 US-09-997-542-239 Sequency 228 18.2 0.7 46 9 US-09-990-427-239 Sequency 229 18.2 0.7 46 9 US-09-990-439-239 Sequency 230 18.2 0.7 46 9 US-09-983-2839 Sequency 231 18.2 0.7 46 9 US-09-983-583-239 Sequency 232 18.2 0.7 46 9 US-09-991-5239 Sequency 233 18.2 0.7 46 9 US-09-992-239 Sequency 234 18.2 0.7 46 9 US-09-992-339 Sequency 235 18.2 0.7 46 9 US-09-992-333-239 Sequency 236 18.2 0.7 46 9 US-09-993-333-239 Sequency 237 18.2 0.7 46 10 US-09-989-722-239 Sequency 238 18.2 0.7 46 10 US-09-989-723-239 Sequency 239 180-2 0.7 46 10 US-09-989-723-239 Sequency 230 180-2 0.7 46 10 US-09-989-723-239 Sequency 231 180-2 0.7 46 10 US-09-989-723-239 Sequency 232 180-2 0.7 46 10 US-09-989-723-239 Sequency 233 180-2 0.7 46 10 US-09-989-723-239 Sequency 234 180-2 0.7 46 10 US-09-989-723-239 Sequency 235 180-2 0.7 46 10 US-09-989-723-239 Sequency 236 180-2 0.7 46 10 US-09-989-723-239 Sequency 237 180-2 0.7 46 10 US-09-989-723-239 Sequency 238 180-2 0.7 46 10 US-09-989-723-239 Sequency 239 180-2 0.7 46 10 US-09-989-723-239 Sequency 239 180-2 0.7 46 10 US-09-989-723-239 Sequency 230 180-2 0.7 46 10 US-09-989-723-239 Sequency   | 210 18.2 0.7 46 9 US-09-97-601-239 Sequenc<br>218 18.2 0.7 46 9 US-09-98-729A-239 Sequenc<br>219 18.2 0.7 46 9 US-09-980-729A-239 Sequenc<br>210 18.2 0.7 46 9 US-09-991-854-239 Sequenc<br>221 18.2 0.7 46 9 US-09-991-854-239 Sequenc<br>221 18.2 0.7 46 9 US-09-997-40-239 Sequenc<br>222 18.2 0.7 46 9 US-09-997-40-239 Sequenc<br>223 18.2 0.7 46 9 US-09-997-628-239 Sequenc<br>224 18.2 0.7 46 9 US-09-997-628-239 Sequenc<br>225 18.2 0.7 46 9 US-09-997-683-239 Sequenc<br>226 18.2 0.7 46 9 US-09-993-469-239 Sequenc<br>227 18.2 0.7 46 9 US-09-993-683-239 Sequenc<br>228 18.2 0.7 46 9 US-09-993-683-239 Sequenc<br>229 18.2 0.7 46 9 US-09-993-683-239 Sequenc   | 206 18.2 0.7 46 9 US-09-990-711-239 Sequence 18.2 0.7 46 9 US-09-980-725-239 Sequence 208 18.2 0.7 46 9 US-09-980-156-239 Sequence 208 18.2 0.7 46 9 US-09-990-156-239 Sequence 209 18.2 0.7 46 9 US-09-991-157-239 Sequence 201 18.2 0.7 46 9 US-09-991-172-239 Sequence 201 18.2 0.7 46 9 US-09-991-172-239 Sequence 201 18.2 0.7 46 9 US-09-997-514-239 Sequence 201 18.2 0.7 46 9 US-09-997-573-239 Sequence 201 18.2 0.7 46 9 US-09-990-726-239 Sequence 201 18.2 0.7 46 9 US-09-9 | 186         18.2         0.7         25         10         US-09-866-108-14629         Sequence           187         18.2         0.7         34         10         US-09-94-420-2         Sequence           188         18.2         0.7         43         10         US-09-96-008A-6         Sequence           189         18.2         0.7         45         10         US-10-082-032-2         Sequence           190         18.2         0.7         45         10         US-09-989-298-239         Sequence           191         18.2         0.7         46         9         US-09-989-293A-239         Sequence           192         18.2         0.7         46         9         US-09-989-734-239         Sequence           193         18.2         0.7         46         9         US-09-989-734-239         Sequence           194         18.2         0.7         46         9         US-09-989-730-239         Sequence           195         18.2         0.7         46         9         US-09-989-730-239         Sequence           196         18.2         0.7         46         9         US-09-991-181-239         Sequence           198  | 171 18.6 0.7 46 10 US-09-738-847-21 Sequence 172 18.6 0.7 50 9 US-09-754-066-15 Sequence 173 18.6 0.7 51 10 US-09-826-630-7 Sequence 174 18.6 0.7 51 10 US-09-826-630-8 Sequence 175 18.6 0.7 51 US-09-826-630-8 Sequence 175 18.6 0.7 55 9 US-09-911-209-72 Sequence 176 18.6 0.7 55 9 US-09-911-209-72 Sequence 177 18.4 0.7 55 9 US-09-911-296A-5140 Sequence 178 18.4 0.7 31 9 US-09-730-289B-2508 Sequence 179 18.4 0.7 38 9 US-09-730-289B-2508 Sequence 179 18.4 0.7 38 9 US-09-730-289B-2508 Sequence 180 18.4 0.7 45 9 US-10-209-952-5 Sequence 181 18.4 0.7 59 10 US-09-870-39A-6616 Sequence 181 18.4 0.7 59 10 US-09-870-39A-6616 Sequence 181 18.4 0.7 59 10 US-09-870-39A-6616 Sequence 181 18.4 0.7 60 9 US-09-90-9941-2001 Sequence 184 18.4 0.7 60 9 US-09-902-941-2001 Sequence 184 18.4 0.7 60 9 US-09-902-941-2001 Sequence 185 186 186 0.7 60 9 US-09-902-941-2001 Sequence 185 186 0.7 60 9 US-09-902-941-2001 Sequence 186 0.7 | 166 18.8 0.7 49 9 US-09-151-771-6 Sequence 167 18.8 0.7 51 10 US-09-808-338-45 Sequence 168 18.6 0.7 25 10 US-09-866-108-14626 Sequence 169 18.6 0.7 38 9 US-09-930-423-2600 Sequence 170 18.6 0.7 42 9 US-10-223-074-70 Sequence |

| 304<br>305<br>306<br>307<br>310  |  | 296<br>297<br>c 298<br>c 299  | 293<br>c 294<br>295   | 290<br>291<br>c 292   |  | c 285                                  |   |   |  | 274<br>c 275<br>c 276  | 2/1<br>c 272<br>273   |  | 266<br>267<br>268   | c 264<br>c 265                         | c 262<br>263                            | 260<br>261                              | .c 258                                  | 255<br>256                             | 253<br>254                               | 251<br>252                               | 249<br>250                           |  |  |  | C 241<br>C 242  |  |
|--|--|---|---|---|--|--|---|---|--|--|---|--|---|--|---|---|---|--|--|--|--------------------------------------|--|--|--|---|--|
| 17.88<br>17.88<br>17.88<br>17.88   | 7777   | 7777  |   | 17.8<br>17.8<br>17.8  |  |  | 17.8<br>17.8  |   | $\perp$                                | 18<br>18   | 18<br>18  | 18<br>18                                   | 18<br>18  | 18                                     |   |   |   | 18.2                                   |  |  |                                      |  |  |  |   |  |
|  |  |   |   |   |  |  |   |   |  |  |   |  |   |  |   |   |   |  |  |  |                                      |  |  |  |   |  |
| 0000005555<br>4400000555555555555555555555   |  | •   |   |   |  |  |   |   |  |  |   |  |   |  |   |   |   |  |  |  |                                      |  |  |  |   |  |
| 10001  | ••••   | 901   | 10  | 999   | 9 1  | 999                                    | 10  | 900   | 999                                    | 979  | 0.0   | 999  | 900   | 999                                    | 9 9                                     | 7 (                                     | 970                                     | 900                                    | 99                                       | 9 9                                      | 9 0                                  | 10<br>10                               | 10                                     | 10                                     | 10  | 10                                     |
| US-10-041-414-31<br>US-10-033-996-29<br>US-9-878-574-1776<br>US-10-057-408-37<br>US-99-735-705-370<br>US-99-850-716A-370<br>US-99-897-778-370<br>US-99-866-108-14655   | US-10-287-919-1219<br>US-09-907-111-267<br>US-09-907-111-269<br>US-10-004-427-47 | US-10-073-300-14<br>US-10-073-300-15<br>US-09-859-104-4<br>US-10-004-427-11 | US-10-152-297-17<br>US-09-790-417-117<br>US-09-790-417-118  | US-10-199-330-14<br>US-10-199-334-14<br>US-10-152-297-16    | US-09-909-496-9<br>US-09-780-533A-6395<br>US-09-877-478-5639 | US-09-465-925-4<br>US-10-269-229-9     | US-10-150-051-4<br>US-09-943-692-5<br>US-10-103-003-3 | JS-09-912-263-503<br>US-09-801-274-1403<br>JS-09-930-423-2953 | JS-10-059-271-60<br>JS-09-888-413-26   | JS-09-918-156-60<br>JS-08-781-986A-3448<br>JS-09-907-111-256 | US-09-930-423-4188<br>US-09-952-060-18<br>US-09-753-436-72  | US-09-848-754A-8144<br>US-09-848-754A-8449 | JS-10-081-408-7<br>JS-10-057-408-42<br>JS-10-076-631-4      | JS-09-915-043-42<br>JS-09-828-062-30   | JS-09-907-111-277<br>JS-09-741-744A-143 | JS-10-055-713-25<br>JS-08-781-986A-5156 | JS-08-781-986A-1748<br>JS-10-085-056-37 | JS-10-138-905-68<br>JS-10-138-916-68   | JS-09-877-478-5801<br>JS-09-877-478-5849 | JS-09-780-533A-5788<br>JS-09-849-928-194 | JS-10-138-838-68<br>JS-10-139-031-68 | US-09-989-721-239<br>JS-10-066-960-194 | US-09-993-604-239<br>US-09-990-456-239 | US-09-990-442-239<br>US-09-991-163-239 | US-09-989-732-239<br>US-09-991-073-239<br>US-09-990-443-239 | US-09-989-727-239<br>US-09-989-731-239 |
| Sequence 31, Appl<br>Sequence 29, Appl<br>Sequence 1776, Appl<br>Sequence 37, Appl<br>Sequence 370, App<br>Sequence 370, App<br>Sequence 370, App  | ~1 m m k1 l  |   |   | 440   | 639  | Sequence 4, Appli<br>Sequence 9, Appli | م.<br>س   | ນີ້ ຫ   | 200                                    | Sequence 60, Appl<br>Sequence 3448, Ap<br>Sequence 256, App  | 12.4  | equence 8                                  | 4 4 7   | uence 4;                               | equence 27                              | equence 25,                             | 174                                     | equence 68<br>equence 68               | equence 580<br>equence 584               | equence 578                              | equence 68,                          | equence 239,<br>quence 194, <i>P</i>   | nce 239,<br>nce 239,                   | equence 239,                           | ence 239,<br>ence 239,                                      | equence 239, equence 239,              |
|  |  |   |   |   |  |  | -   |   |  | ***  |   |  |   |  |   |   |   |  |  |  |                                      |  |  |  |   |  |
| 378<br>0 378<br>0 380<br>0 381<br>0 383  | 373<br>C 374<br>C 375  | 369<br>370<br>371<br>372  | C 366<br>C 367<br>C 368   | C 363<br>364<br>365   | 360<br>361<br>362  | C 358                                  |   | 3 3 3 3<br>3 5 5 5<br>4 8 8                                   | c 350<br>351                           |  | C 344<br>C 345  |  | 339<br>C 340  |  | C 335                                   |   | c 331                                   | C 328                                  | 326<br>327                               | 32 <b>4</b><br>325 .                     | 322<br>323                           |  | c 318<br>c 319                         |  | c 314   |  |
| 3377<br>3777<br>3879<br>3880<br>3883   | 373 17.<br>374 17.<br>375 17.<br>376 17.   | 369 17.4<br>370 17.4<br>371 17.4<br>372 17.4                                | 366 1<br>367 1<br>368 1   |   | 360 17.4<br>361 17.4<br>362 17.4                             |  | 355 17.4<br>356 17.4                                  | 352 17.4<br>353 17.4<br>354 17.4                              |  |  |   | 342 17.<br>343 17.                         |   | 337 17.<br>338 17.                     | 335 17.<br>336 17.                      | 333 17.                                 | 331 17.<br>332 17.                      | 328 17.<br>329 17.                     | 17.<br>17.                               | 17                                       | 17.<br>17.                           | 320 17.<br>321 17.                     | 318 17.<br>319 17.                     | 317 17.                                | 314 17.<br>315 17.  | 312 17.<br>313 17.                     |
| 377 17.<br>378 17.<br>379 17.<br>380 17.<br>381 17.<br>382 17.<br>384 17.  | 373 17.4 0.<br>374 17.4 0.<br>375 17.4 0.<br>376 17.4 0.                         | 17.4 0<br>17.4 0<br>17.4 0<br>17.4 0  | 366 17.4 0.367 17.4 0.368 17.4 0.   | 363 17.4 0.<br>364 17.4 0.<br>365 17.4 0.                   | 17.4 0.<br>17.4 0.<br>17.4 0.                                | 358 17.4 0.<br>359 17.4 0.             | 17.4 0.   | 17.4 0.<br>17.4 0.<br>17.4 0.                                 | 350 17.4 0.<br>351 17.4 0.             | 347 17.4 0.<br>348 17.4 0.<br>349 17.4 0.                    | 344 1/.4 0.<br>345 17.4 0.<br>346 17.4 0.                   | 342 17.6 0.<br>343 17.4 0.                 | 339 17.6 0.<br>340 17.6 0.<br>341 17.6 0.                   | 337 17.6 0.<br>338 17.6 0.             | 335 17.6 0.<br>336 17.6 0.              | 333 17.6 0.<br>334 17.6 0.              | 331 17.6 0.<br>332 17.6 0.              | 328 17.6 0.<br>329 17.6 0.             | 17.6 0.<br>17.6 0.                       | 17.6 0.                                  | 17.6 0.<br>17.6 0.                   | 320 17.6 0.<br>321 17.6 0.             | 318 17.6 0.<br>319 17.6 0.             | 317 17.6 0.                            | 314 17.6<br>315 17.6  | 31.2 17.6 0.<br>31.3 17.6 0.           |
| 377 17.4 0.3 378 17.2 0.3 379 17.2 0.3 380 17.2 0.3 382 17.2 0.3 383 17.2 0.3 384 18.2 0.3 384 18.2 0.3 384 18.2 0.3 384 18.2 0.3 384 18.2 0.3 384 1 | 373 17.4 0.6<br>374 17.4 0.6<br>375 17.4 0.6<br>376 17.4 0.6                     | 17.4 0.6<br>17.4 0.6<br>17.4 0.6<br>17.4 0.6                                | 366 17.4 0.6<br>367 17.4 0.6<br>368 17.4 0.6  | 363 17.4 0.6<br>364 17.4 0.6<br>365 17.4 0.6                | 17.4 0.6<br>17.4 0.6<br>17.4 0.6                             | 358 17.4 0.6<br>359 17.4 0.6           | 17.4 0.6  | 17.4 0.6<br>17.4 0.6<br>17.4 0.6                              | 350 17.4 0.6<br>351 17.4 0.6           | 347 17.4 0.6<br>348 17.4 0.6<br>349 17.4 0.6                 | 344 1/.4 0.6<br>345 17.4 0.6<br>346 17.4 0.6                | 342 17.6 0.6<br>343 17.4 0.6               | 339 17.6 0.6<br>340 17.6 0.6<br>341 17.6 0.6                | 337 17.6 0.6<br>338 17.6 0.6           | 335 17.6 0.6<br>336 17.6 0.6            | 333 17.6 0.6<br>334 17.6 0.6            | 331 17.6 0.6<br>332 17.6 0.6            | 328 17.6 0.6<br>329 17.6 0.6           | 17.6 0.6<br>17.6 0.6                     | 17.6 0.6                                 | 17.6 0.6<br>17.6 0.6                 | 320 17.6 0.6<br>321 17.6 0.6           | 318 17.6 0.6<br>319 17.6 0.6           | 317 17.6 0.6                           | 314 17.6 0.6<br>315 17.6 0.6                                | 313 17.6 0.6                           |
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| 10-09-79 10-09-79 10-09-79 10-09-79 10-09-79 10-09-79 10-09-79 10-09-79 10-12  | 5-963-24<br>-1653-29<br>-5-543-3092 ·<br>-135-19<br>-135-19<br>-11-843-11<br>-843-11   |
| 0.9-796-692-9235 1.0-040-662-9235 1.0-125-648-10 0.9-362-286-27 1.0-03-22-86-32 1.0-007-280A-2 1.0-013-737-11 0.9-954-692-26 1.0-228-070-11 1.0-125-635A-357 0.9-938-864-357 0.9-938-864-357 10-022-603-357 10-022-603-357 10-120-687-19 10-136-891-17 10-120-687-19 10-9801-274-36 -09-991-980-3 10-984-754-3907 -09-884-754A-3907 -09-884-754A-3907 -09-874-470-11 0-12-488-61 10-112-488-61 10-112-488-61 10-112-38-28 09-991-3-28-28 09-991-3-28-28 09-991-3-28-28 09-991-3-28-28 09-991-3-28-28 09-913-238-28 09-913-238-28 09-913-238-28 09-913-238-28 09-913-238-28 09-913-238-28 09-913-238-28 09-913-238-28 09-913-238-28 09-913-238-28 09-913-238-28   | •  |
| 0.9-796-692-9235 1.0-040-662-9235 1.0-125-648-10 0.9-362-286-27 1.0-03-22-86-32 1.0-007-280A-2 1.0-013-737-11 0.9-954-692-26 1.0-228-070-11 1.0-125-635A-357 0.9-938-864-357 0.9-938-864-357 10-022-603-357 10-022-603-357 10-120-687-19 10-136-891-17 10-120-687-19 10-9801-274-36 -09-991-980-3 10-984-754-3907 -09-884-754A-3907 -09-884-754A-3907 -09-874-470-11 0-12-488-61 10-112-488-61 10-112-488-61 10-112-38-28 09-991-3-28-28 09-991-3-28-28 09-991-3-28-28 09-991-3-28-28 09-991-3-28-28 09-913-238-28 09-913-238-28 09-913-238-28 09-913-238-28 09-913-238-28 09-913-238-28 09-913-238-28 09-913-238-28 09-913-238-28 09-913-238-28 09-913-238-28   | •  |
| 09-796-692-9235 10-040-682-9235 10-025-648-10 09-362-286-27 10-036-286-32 10-007-2808-32 10-007-2808-2 110-013-737-11 09-954-692-26 10-28-070-11 10-125-631A-357 09-955-9671A-26 110-202-603-357 10-002-603-357 10-126-687-19 10-136-891-17 10-120-687-19 10-136-891-17 10-120-687-19 10-991-9801-274-353 -09-991-9801-274-353 -09-991-9801-274-353 -09-911-8-110 -09-874-470-11 -09-874-470-12 -10-108-714-5 09-975-237-238-29 09-913-238-27 09-913-238-27 09-913-238-27 09-913-238-27 09-913-238-28 10-045-27-353 09-913-238-28 09-913-238-28 09-913-238-28 09-913-238-28 09-913-238-28  | •  |
| 09-796-692-9235 10-040-682-9235 10-0125-648-10 09-362-286-27 10-036-286-32 10-007-280A-2 10-007-280A-2 11-013-737-11 09-955-9-671A-26 10-025-671A-26 10-025-635A-357 09-963-875-19 10-126-687-19 -09-801-274-353 -09-991-980-317-09-801-274-353 -09-991-980-374-3907 -09-801-274-353 -09-913-238-310 09-975-272-353 09-913-238-28 110-012-538-21 09-975-272-353 09-913-238-28 09-975-272-353 09-913-238-28 09-975-23-28 09-913-238-28 110-012-38-28 09-975-23-28 09-913-238-28 09-975-23-28 09-913-238-28 09-975-23-28 09-913-238-28 09-975-23-28 09-913-238-28 09-975-23-28 09-913-238-28 09-975-23-28 09-913-238-28  | •  |
| 0.9-796-692-9235 10-10-125-648-10 10-125-648-10 0.9-362-286-32 10-007-2806-32 10-007-2806-32 10-013-737-11 09-954-692-26 09-995-529-118 3:09-559-671A-26 11-125-635A-357 09-981-864-357 09-981-891-17 10-120-687-19 10-120-687-19 10-98-11-274-203 -09-91-91-980-3 10-056-414-735 -09-91-81-17 10-120-687-19 10-120-687-19 10-120-687-19 10-120-687-19 09-91-288-61 10-043-573-110 09-913-238-27 09-913-238-27 09-913-238-28 09-913-238-28 09-913-238-28 09-913-238-28 09-913-238-28 09-913-238-28 09-913-238-28 09-913-238-28   | •  |

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Query Match
Best Local Similarity
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: Sequence 45, Application US/10057408 

: Publication No. US20030082561A1
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US-10-057-408-49
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RESULT 4
US-10-057-408-40
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Publication No. US20030082561A1
GENERAL INFORMATION:
APPLICANT: Sera, Takashi
TITLE OF INVENTION: Zinc Finger Domain Recognition Code and Uses Thereof
FILE REFERENCE: 109845.135
CURRENT APPLICATION NUMBER: US/10/057,408
CURRENT FILING DATE: 2002-01-23
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Best Local
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LENGTH: 56
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Best Local
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TITLE OF INVENTION: Zinc Finger Domain Recognition Code and Uses
FILE REFERENCE: 109845.135
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PRIOR APPLICATION NUMBER: US 60/220,060
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SOFTWARE: PatentIn version 3.0
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                                                                                                                                                                                                                                              ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                TYPE: DNA
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les 35; Conserv
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Pred. No. 3.3e
0; Mismatches
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Pred. No. 1.3e+02;
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Pred. No. 1.7e+02;
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RESULT 6
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LENGTH: 56
TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                             GENERAL INFORMATION:
                                                                                                                                              Sequence 50, Appropriate Publication No.
                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
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APPLICANT: Sera, Takashi
TITLE OF INVENTION: Zinc Finger Domain Recognition Code and Uses Thereof
ETLE REFERENCE: 109845.135
CURRENT APPLICATION NUMBER: US/10/057,408
CURRENT FILING DATE: 2002-01-23
PRIOR APPLICATION NUMBER: US 60/220,060
PRIOR FILING DATE: 2000-07-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: C. Frank Bennett
APPLICANT: Susan M. Freier
TITLE OF INVENTION: ANTISENSE MODULATION OF
FILE REFERENCE: RTS-0248
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/10/057,408
CURRENT FILING DATE: 2002-01-23
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION NUMBER: US/09/898,556A CURRENT FILING DATE: 2001-07-03 .
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SOFTWARE: PatentIn version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OTHER INFORMATION: Nucleotides 48-50 and 54-58 are "n" wherein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       OTHER INFORMATION: Partial zinc finger domain oligomer NAME/KEY: misc_feature LOCATION: (48)...(58)
                                                                                                                                                                                                                                                                                                                                                                                                                                     FEATURE:
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31; Conser
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0; Mismatches
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Pred. No. 3.
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RESULT 8
US-10-057-408-41/c
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                                                                                      SEQ ID NO 41
LENGTH: 55
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Best Local
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Publication No. US20030036641A1
                                                                                                                                PRIOR APPLICATION NUMBER: US 60/220,060 PRIOR FILING DATE: 2000-07-21 NUMBER OF SEQ ID NOS: 69
                                                                                                                                                                                        CURRENT APPLICATION NUMBER: US/10/057,408 CURRENT FILING DATE: 2002-01-23
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                                                                                                                                                                                                                          TITLE OF INVENTION: Zinc Finger Domain Recognition Code and Uses Thereof FILE REFERENCE: 109845.135
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                                                                                                                       SOFTWARE: PatentIn version 3.0
                                                                                                                                                                                                                                                           APPLICANT: Sera,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION NUMBER: US/09/775,049
CURRENT FILING DATE: 2001-01-31
NUMBER OF SEQ ID NOS: 40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Lindbo, John A.
TITLE OF INVENTION: Methods For Homology-Driven Reassembly
TITLE OF INVENTION: of Nucleic Acid Sequences
FILE REFERENCE: P-LG 4412
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Fitzmaurice, APPLICANT: Lindbo, John
              OTHER INFORMATION: Partial zinc finger domain oligomer
                                                 ORGANISM: Artificial Sequence
                                                                      TYPE: DNA
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SOFTWARE: PatentIn version 3.0
NAME/KEY: misc_feature
                                    FEATURE:
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Similarity 65.5%;
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Pred. No. 1.3e+03;
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Pred. No. 7.9e+02;
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; OTHER INFORMATION: Description of Artificial Sequence: oligonucleotide PB2
US-10-055-713-24
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US-10-055-713-24
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; OTHER INFORMATION: primer
US-09-906-393A-34
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 Query Match
Best Local Similarity
                                                                                                                                                                     SEQ ID NO 24
                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 24, Application US/10055713 Publication No. US20030044957A1
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Best Local Similarity 56.4
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Best Local Similarity
Matches 32; Conserv
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                                                                                                                                                                                                       PRIOR APPLICATION NUMBER: 60/290,716
PRIOR FILING DATE: 2001-05-11
NUMBER OF SEC. 75
                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: ZINC FINGER PROTEINS FOR DNA BINDING AND GENE TITLE OF INVENTION: REGULATION IN PLANTS
FILE REFERENCE: 8325-0026 / S26-US1
CURRENT APPLICATION NUMBER: US/10/055,713
CURRENT FILING DATE: 2002-06-17
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CURRENT FILING DATE: 2001-07-16
PRIOR APPLICATION NUMBER: 60/218,761
PRIOR FILING DATE: 2000-07-17
                                                                                                        LENGTH: 54
TYPE: DNA
ORGANISM: Artificial
                                                                                                                                                                                  SOFTWARE:
                                                                                                                                                                                    NUMBER OF SEQ ID NOS: 105
SOFTWARE: PatentIn Ver. 2.0
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TITLE OF INVENTION: METHOD OF PROGNOSING CANCER AND
FILE REFERENCE: 1720-1-001CIP
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SOFTWARE: PatentIn version 3.1
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                                                                                           FEATURE:
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ORGANISM: Artificial sequence
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No. US20030039970A1
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72.5%;
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56.48;
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Score 22.4; DB 9;
Pred. No. 4.2e+03;
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Pred. No.
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Pred. No. 1.5e+03;
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2.9e+03;
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                                                       Db
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US-10-057-408-48/c
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RESULT 13
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Best Local
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LENGTH: 60
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LENGTH: 60
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CURRENT FILING DATE: 2002-01-23
PRIOR APPLICATION NUMBER: US 60/220,060
PRIOR FILING DATE: 2000-07-21
NUMBER OF SEQ ID NOS: 69
                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: Zinc Finger Domain Recognition Code and Uses Thereof FILE REFERENCE: 109845.135
CURRENT APPLICATION NUMBER: US/10/057,408
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TITLE OF INVENTION: Zinc Finger Domain Recognition Code and
FILE REFERENCE: 109845.135
                                                                                                                                                                                                                                                                                                                                                                 PRIOR FILING DATE: 2000-07-21 NUMBER OF SEQ ID NOS: 69
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Sera, Takashi
                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION NUMBER: US 60/220,060
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                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: PatentIn version 3.0
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OTHER INFORMATION: Nucleotides 37-39 and 46-51 are "n" wherein "n"
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ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                                                           TYPE: DNA
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                                                                                                                                                   Local Similarity
                                                                                           1239 CACCAGAGGCGCACACTGGGGAGAAGCCTTA 1270
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                                                       36 CATCAGCGTACTCACACTGGCGAGAAGCCTTA 5
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Pred. No. 4
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Pred. No. 4.5e+03;
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                                                                                                                                                                     US-09-898-556A-4
                                                                                                                                                                                         RESULT 15
                                                                                                         Sequence 4, Application US/09898556A Publication No. US20030087849A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 33, Application US/09906393A Publication No. US20030039970A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQ ID NO 33
LENGTH: 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 182
                                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Publication No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 182, Application US/09912263
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Ireland, James S.
APPLICANT: Lander, Eric. S.
TITLE OF INVENTION: HUMAN SINGLE NUCLEOTIDE POLYMORPHISMS
FILE REFERENCE: 2825.2017-001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/09/906,393A CURRENT FILING DATE: 2001-07-16 PRIOR APPLICATION NUMBER: 60/218,761 PRIOR FILING DATE: 2000-07-17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Wang, Zhou
APPLICANT: Xiao, Wuhan
TITLE OF INVENTION: METHOD OF PROGNOSING CANCER AND THE PROTEINS INVOLVED
FILE REFERENCE: 1720-1-001CIP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION NUMBER: US/09/912,263
CURRENT FILING DATE: 2001-07-24
                                                     APPLICANT: Susan M. Freier TITLE OF INVENTION: ANTISENSE MODULATION OF HKR1
                                                                         APPLICANT: C. Frank Bennett APPLICANT: Susan M. Freier
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQ ID NOS: 552
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR FILING DATE: 2000-07-24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION NUMBER: US 60/220,315
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Cargill, Michele
CURRENT APPLICATION NUMBER: US/09/898,556A CURRENT FILING DATE: 2001-07-03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: PatentIn version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQ ID NOS: 36
                                   FILE REFERENCE: RTS-0248
                                                                                                                                                                                                                                                                                                                                                                                                                     OTHER INFORMATION: primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                           ORGANISM: Artificial sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                             FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH: 31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1224 TCAAACCTCATTACCCACCAGAGGGCGCA 1252
                                                                                                                                                                                                                                                                 89 TCAGGATCCGCGGGGTGCACCCGCGTTCCATCTGTGTCTTCTGAGACTT 135
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   24;
                                                                                                                                                                                                                                                                                                                             31;
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                                                                                                                                                                                                                                               6 TCTGGATCCGAGGTGAGTACTGACTTTACACACGTCTTCCGAAGCTT
                                                                                                                                                                                                                                                                                                                                             Similarity
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                                                                                                                                                                                                                                                                                                                     Score 21.4; DB y; Pred. No. 9.8e+03;
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Pred. No. 3.4e+03;
1; Mismatches 4
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Length 31;

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Gaps

0,

EXPRESSION

Length 60; Indels

0

Gaps

0;

SEQ ID NO 4

NUMBER OF SEQ

IJ

NOS:

89

Length 45

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RESULT 17
US-10-057-408-56/c
US-10-057-408-56/c
; Sequence 56, Application US/10057408
; Publication No. US20030082561a1
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                             В
; OTHER INFORMATION: PCR primer US-10-057-408-56
                                                                                                                                                                                                                                                                                                                                                                                                                           δÃ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-801-274-1402
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                                                                                               SOFTWARE: PatentIn version 3.0 SEQ ID NO 56
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local (
                                                                                                                                  PRIOR APPLICATION NUMBER: US 60/220,060 PRIOR FILING DATE: 2000-07-21 NUMBER OF SEQ ID NOS: 69
                                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/10/057,408
CURRENT FILING DATE: 2002-01-23
                                                                                                                                                                                                                    TITLE OF INVENTION: Zinc Finger Domain Recognition Code and Uses Thereof FILE REFERENCE: 109845.135
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Cargill, Michele
APPLICANT: Ireland, James S.
APPLICANT: Lander, Eric S.
TITLE OF INVENTION: HUMAN SINGLE NUCLEOTIDE POLYMORPHISMS
FILE REFERENCE: 2825.2009-001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION NUMBER: US 60/187,510
PRIOR FILING DATE: 2000-03-07
PRIOR APPLICATION NUMBER: US 60/206,129
PRIOR FILING DATE: 2000-05-22
                               FEATURE:
                                            ORGANISM: Artificial Sequence
                                                                 LENGTH: 45
TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: FastSEQ for Windows Version 4.0
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CURRENT FILING DATE: 2001-03-07
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TYPE: DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local Similarity les 24; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                      1358 TTGCAGGGAGTGTGAGCAAGGCTTTAGCCAG 1388
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2276 TGAAGGAGAATTGCTGGCTCA 2296
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                                                                                                                                                                                                                                                                                                                                                                                          TTGCAAGGAGTGTGGRAAGGCCTTCAGCCAG 31
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 21; DB 10;
Pred. No. 8.6e+03;
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; SEQUENCE DESCRIPTION: SEQ ID NO: 236: US-10-066-960-236
Query Match
Best Local Similarity
Watches 28; Conserva
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 236, Application US/10066960 Publication No. US20030049644A1 GENERAL INFORMATION:
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Best Local:
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                                                                                                                                                                                                                                                                             TELEFAX: (303) 793-34 INFORMATION FOR SEQ ID NO: 236: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local Similarity
                                                                                                                                          FEATURE:
                                                                                                                                                                            FEATURE:
                                                                                                                                                                                            MOLECULE TYPE: RNA
                                                                                                                                                                                                                                                                                                                                                 TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                      ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3 1/2 diskette, 1.44 MB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION DATA:
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ADDRESSEE: Swanson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: HIGH AFFINITY NUCLEIC ACID LIGANDS TO LECTINS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1239 CACCAGAGGGCGCACACTGGGGAGAAGCC 1267
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: PARMA, et al
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    45 CACCAACGCACGCACACGGGGAGAAGCC 17
                                                                                                                                                                                                                           TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US/10/066,960 FILING DATE: 04-Feb-2002 CLASSIFICATION: (Unknown)
                                                                                                                                                                                                            TOPOLOGY:
                                                                                                                                                                                                                                                                                                                              TELEPHONE: (303)
                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE: 07-JUNE-1995
APPLICATION NUMBER: 08/477,829
FILING DATE: 07-JUNE-1995
                                                                                                                                                        OTHER INFORMATION:
                                                                                                                                                                                                                                                             LENGTH: 59 base pairs
                                                                                                                                                                                                                                                                                                                                                                REGISTRATION NUMBER: 33,215
REFERENCE/DOCKET NUMBER: NEX40C/PCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM pc compatible OPERATING SYSTEM: MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: 08/472,255
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: PCT/US96/09455
FILING DATE: 05-JUNE-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE: 1999-DEC-03
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STREET: 8400 E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: 08/472,256 FILING DATE: 07-JUNE-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: 08/479 FILING DATE: 07-JUNE-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: 08/952,793
                 Conservative
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                                                                                                                                                                                                           linear
                                0.8%;
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                Score 21; DB
Pred. No. 1.3e
2; Mismatches
                                                                                                                                                      All C's are 2'-F cytosine
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Pred. No. 1.1e+04;
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                                1.3e+04;
                                                DB 9; Length 59;
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                Indels
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                Gaps
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QΥ

140 CTTCTCCAGGAAGAGCACTCAGGAGACCAGGAAAATGGCTACAGG 184

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RESULT 19
US-09-849-928-236
; Sequence 236, App.
; Publication No. U
                                                                                                                                                                                                                  SEQUENCE DESCRIPTION: All U's are 2'-F uracil US-09-849-928-236
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                  RESULT 20
US-10-066-960-244
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                                                                                                                  QΥ
; Sequence 244, Application US/10066960
                                                                                                                                                                                           Query Match
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                                                                                                                                                        Local Similarity
les 28; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/849,928
FILING DATE: 04-May-2001
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM: MEDIUM TYPE: Diskette, 3 1/2 diskette, 1.44
                                                                                                                                                                                                                                                                                FEATURE:
                                                                                                                                                                                                                                                                                                                  FEATURE:
                                                                                                                                                                                                                                                                                                                                   MOLECULE TYPE: RNA
                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEPHONE: (303) 793-3333
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT:
                                                                                                                    140 CTTCTCCAGGAAGAGCACTCAGGAGACCAGGAAAATGGCTACAGG 184
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: 08/479,724
FILING DATE: 07-JUNE-1995
APPLICATION NUMBER: 08/472,256
FILING DATE: 07-JUNE-1995
APPLICATION NUMBER: 08/472,255
FILING DATE: 07-JUNE-1995
APPLICATION NUMBER: 08/477,829
FILING DATE: 07-JUNE-1995
FILING DATE: 07-JUNE-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM pc compatible OPERATING SYSTEM: MS-DOS SOFTWARE: WordPerfect 6.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CUCAACGAGCAAAAGUACUCACGGGACCAGGAGAUCAGCAACACG 45
                                                                                     STATE: Colorado
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CITY: Englewood
                                                                                                                                                                                                                                                                                                                                                                              TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY: USA
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                                                                                                                                                                                                                                                                                                OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                    STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEFAX: (303)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              REFERENCE/DOCKET NUMBER: NEX40C/PCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: Barry J. Swanson REGISTRATION NUMBER: 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: 08/952,793 FILING DATE: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Application US/09849928
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           INVENTION: HIGH AFFINITY NUCLEIC ACID LIGANDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US20030059769A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        8400 E. Prentice Avenue, Suite 200
                                                                                                                                                                                                                                                                                                                                                                                                                                         (303) />
(303) />
(303) 793-3433
                                                                                                                                                                                                                                                                                                                                                     linear
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                                                                                                                                                                        0.8%;
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                                                                                                                                                                       Score 21;
Pred. No.
                                                                                                                                                                                                                                                                                               All C's are 2'-F cytosine
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                                                                                                                                                        Mismatches
                                                                                                                                                      nes 15;
                                                                                                                                                                                           DB 9;
                                                                                                                                                                                       Length 59;
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                                                                                                US-09-849-928-244
                                                                                                                   RESULT 21
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                       Sequence 244, Application US/US
Publication NO. US20030059769A1
GENERAL INFORMATION:
APPLICANT: PARMA, et al.
                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Publication No. US20030049644A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX: (303) 79 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION DATA:
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MEDIUM TYPE: Diskette,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: PARMA, et al. TITLE OF INVENTION: HIGH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE: 07-JUNE-1995 ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
            TITLE OF INVENTION: HIGH AFFINITY NUCLEIC ACID LIGANDS
                                                                                                                                                                                                                                                                                                                              SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                 FEATURE
                                                                                                                                                                                                                                                                                                                                                                                                                   MOLECULE TYPE: RNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                   FEATURE
                                                                                                                                                                                      580 CAGAAGATCAGAAACAACAGCAGGATCCATTCTGC 614
                                                                                                                                                                      4 CAAGCGGUCAGAAACAAUAGCUGGAUACAUACCGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 52 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US/10/066,960 FILING DATE: 04-Feb-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER: IBM pc compatible OPERATING SYSTEM: MS-DOS
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                                                                                                                                                                                                                                                                                                                                               OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                               OTHER INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: 08/952,793
FILING DATE: 1999-DEC-03
APPLICATION NUMBER: PCT/US96/09455
FILING DATE: 05-JUNE-1995
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEPHONE: (303) 793-3333
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: 08/472,255 FILING DATE: 07-JUNE-1995
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: 08/477,829
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PPLICATION NUMBER: 08/472,256
                                                                               Application US/09849928
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LING DATE:
                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                              DESCRIPTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Barry J. Swanson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     EE: Swanson & Bratschun, L.L.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    USA
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                                                                                                                                                                                                                                                         0.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          HIGH AFFINITY NUCLEIC ACID LIGANDS
TO LECTINS
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                                                                                                                                                                                                                                                                                                                              SEQ
                                                                                                                                                                                                                                                        Score 20.6;
Pred. No. 1
                                                                                                                                                                                                                                                                                                                             All U's are 2'-F uracil
Q ID NO: 244:
                                                                                                                                                                                                                                                                                                                                                                               All C's
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              33,215
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                                                                                                                                                                                                                                           Mismatches
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                                                                                                                                                                                                                                                                                                                                                                               are 2'-F cytosine
                                                                                                                                                                                                                                                           .7e+04;
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US-09-950-442-40/c
                                                                                                                                                                                                                                                                                                                                     В
                                                                                                                                                                                                   Sequence 40, Application US/09950442 Publication No. US20030027127A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                      Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX: (303) 793-:
INFORMATION FOR SEQ ID NO: 244
SEQUENCE CHARACTERISTICS
                                                     CORRESPONDENCE ADDRESS:
ADDRESSEE: James F. Haley,
STREET: 1251 Avenue of the
                                                                                                                             APPLICANT: Farr, Spencer B.

TITLE OF INVENTION METHODS AND DIAGNOSTIC

DETERMINING TOXICITY
                                                                                                              NUMBER OF SEQUENCES:
                                                                                                                                                                    APPLICANT: Farr, Todd,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     REFERENCE/DOCKET NUMBER: NEX40C/PCT TELEPHONE: (303) 793-3333
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ZIP: 80111
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MOLECULE TYPE: RNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                             580 CAGAAGATCAGAAACAACAGCAGGATCCATTCTGC 614
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQUENCES: 390 CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                       23;
                                                                                                                                                                                                                                                                                                                          4 CAAGCGGUCAGAAACAAUAGCUGGAUACAUACCGC 38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: 08/952,793
FILING DATE: <Unknown>
APPLICATION NUMBER: 08/479,724
FILING DATE: 07-JUNE-1995
APPLICATION NUMBER: 08/472,256
FILING DATE: 07-JUNE-1995
APPLICATION NUMBER: 08/472,255
APPLICATION NUMBER: 08/472,255
FILING DATE: 07-JUNE-1995
APPLICATION NUMBER: 08/477,829
FILING DATE: 07-JUNE-1995
APPLICATION NUMBER: 08/477,829
FILING DATE: 07-JUNE-1995
                                        CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US/09/849,928
FILING DATE: 04-May-2001
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM pc compatible OPERATING SYSTEM: MS-DOS SOFTWARE: WordPerfect 6.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LENGTH: 52 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CITY: Englewood
STATE: Colorado
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADDRESSEE: Swanson & Bratschun, L.L. STREET: 8400 E. Prentice Avenue, Sui
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    linear
                                                                                                                                                                                                                                                                                                                                                                                                                    0.78;
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                                                                                                                                                                                                                                                                                                                                                                                                                   Score 20.6;
Pred. No. 1.
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                                                                        Jr. c/o FISH & NEAVE
                                                                                                                                                                                                                                                                                                                                                                                                               1.7e+04;
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                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 9;
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                                                                                                                                            KITS
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                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 52;
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                                                                                                                                                                                                                                                                                                                                                                                            Gaps
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RESULT 24
US-09-898-556A-13/c
; Sequence 13, Application US/09898556A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; SEQUENCE DESCRIPTION: SEQ ID NO: 40: US-09-950-442-40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity 68.3
                                                                                                                                                                                                                              NUMBER OF SEQ ID NOS: 89 SEQ ID NO 12
                                                                   Matches
                                                                                                                                                                                                                                                                                                                                                                      Sequence 12, App
Publication No.
                                                                                  Best Local
                                                                                               Query Match
                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION
                                                                                                                                                                                                                                                    APPLICANT: C. Frank Bennett
APPLICANT: Susan M. Freier
TITLE OF INVENTION: ANTISENSE MODULATION OF I
FILE REFERENCE: RTS-0248
CURRENT APPLICATION NUMBER: US/09/898,556A
CURRENT FILING DATE: 2001-07-03
                                                                                                                                                                 FEATURE:
                                                                                                                                                                           ORGANISM: Artificial Sequence
                                                                                                                                                                                               LENGTH: 20
TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  INFORMATION FOR SEQ ID NO: 40:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2212 TGGAGACATTCTGTGTGTGATTATGCATGAGACTGTACTGG 2252
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TOPOLOGY: linear MOLECULE TYPE: cDNA HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            REGISTRATION NUMBER: 27,794
REFERENCE/DOCKET NUMBER: X-1 CIP
TELECOMMUNICATION INFORMATION:
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COMPUTER READABLE FORM:
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                            33 GCCTCCCTCTTAAGGTCTTT 52
20 GCCTCCCTCTTAAGGTCTTT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   48 TGCAGACATACTGAGTGTCACCGTTGAAGAGAGTGGAGTGG 8
                                                                   20;
                                                                               Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LENGTH: 50 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                          Application US/09898556A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEFAX: (212) 596-90
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: Haley Jr., James F. REGISTRATION NUMBER: 27,7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: 08/374,641 FILING DATE: 12-JUL-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US/09/950,442 FILING DATE: 10-Sep-2001 CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: Patentin Release #1.0,
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                                                                 Conservative
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Pred. No.
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Pred. No. 2.2e+04;
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                                                            1.4e+04;
hes 0;
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US-09-898-556A-14/c
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US-09-898-556A-15/c
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US-09-898-556a-14
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                                                     CURRENT APPLICATION NUMBER: US/09/898,556A
CURRENT FILING DATE: 2001-07-03
NUMBER OF SEQ ID NOS: 89
SEQ ID NO 15
LENGTH: 20
TYPE: DNA
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LENGTH: 20

TYPE: DNA

ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                 Sequence 15, Application US/09898556A Publication No. US20030087849A1 GENERAL INFORMATION:
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LENGTH: 20
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GENERAL INFORMATION
                                                                                                                                                        APPLICANT: C. Frank Bennett
APPLICANT: Susan M. Freier
TITLE OF INVENTION: ANTISENSE MODULATION OF HKR1 EXPRESSION
FILE REFERENCE: RTS-0248
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APPLICANT: Susan M. Freter
TITLE OF INVENTION: ANTISENSE MODULATION OF HKR1 EXPRESSION
FILE REFERENCE: RTS-0248
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APPLICANT: Susan M. Freier
TITLE OF INVENTION: ANTISENSE MODULATION OF HKR1
FILE REFERENCE: RTS-0248
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OTHER INFORMATION: Antisense Oligonucleotide
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RESULT 29
US-09-898-556A-18/c
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; Publication No. US20030087849A1
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; Publication No. US20030087849A1
; GENERAL INFORMATION:
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LENGTH: 20
TYPE: DNA
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Best Local :
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CURRENT APPLICATION NUMBER: US/09/898,556A
CURRENT FILING DATE: 2001-07-03
                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION: ANTISENSE MODULATION OF HKR1 EXPRESSION FILE REFERENCE: RTS-0248
CURRENT APPLICATION NUMBER: US/09/898,556A
CURRENT FILING DATE: 2001-07-03
NUMBER OF SEQ ID NOS: 89
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TYPE: DNA
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                                                                CURRENT APPLICATION NUMBER: US/09/898,556A
CURRENT FILING DATE: 2001-07-03
NUMBER OF SEQ ID NOS: 89
SEQ ID NO 20
LENGTH: 20
                                                                                                                                                                                                               Sequence 20, Application US/09898556A Publication No. US20030087849A1 GENERAL INFORMATION:
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FILE REFERENCE: RTS-0248
CURRENT APPLICATION NUMBER: US/09/898,556A
CURRENT FILING DATE: 2001-07-03
NUMBER OF SEQ ID NOS: 89
SEQ ID NO 19
LENGTH: 20
TYPE: DNA
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LENGTH: 20
TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                         APPLICANT: C. Frank Bennett
APPLICANT: Susan M. Freier
TITLE OF INVENTION: ANTISENSE MODULATION OF HKR1 EXPRESSION
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                         TYPE: DNA
ORGANISM: Artificial Sequence
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EQ ID NO 18
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OTHER INFORMATION: Antisense Oligonucleotide
                      FEATURE:
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nes 20; Conserv
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RESULT 34
US-09-898-556A-23/c
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US-09-898-556A-22/c
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; OTHER INFORMATION: Antisense Oligonucleotide
US-09-898-556A-21
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Sequence 23, Application US/09898556A Publication No. US20030087849A1 GENERAL INFORMATION:
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                                                                                                                                                         Query Match
Best Local Similarity 100.
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SEQ ID NO 22
LENGTH: 20
TYPE: DNA
ORGANISM: Artificial Sequence
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Best Local S
Matches 20
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LENGTH: 20
TYPE: DNA
ORGANISM: Artificial Sequence
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CURRENT APPLICATION NUMBER: US/09/898,556A
CURRENT FILING DATE: 2001-07-03
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Best Local Similarity
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TITLE OF INVENTION: ANTISENSE MODULATION OF HKR1 EXPRESSION
                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: C. Frank Bennett
APPLICANT: Susan M. Freier
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APPLICANT: Susan M. Freier
TITLE OF INVENTION: ANTISENSE MODULATION OF
FILE REFERENCE: RTS-00248
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                                                                                                                                                                                                                                                                   FEATURE:
                                                                                                                       482 TCAGCAAGCTCTCAGCCAAC 501
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Pred.
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0; Mismatches
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NO.
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FEATURE:
OTHER INFORMATION: Antisense Oligonucleotide
US-09-898-556A-23
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US-09-898-556A-24/c
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US-09-898-556A-25/c
Sequence 25, Application US/09898556A
Publication No. US20030087849A1
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OTHER INFORMATION: Antisense Oligonucleotide US-09-898-556A-24
US-09-898-556A-25
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LENGTH: 20
TYPE: DNA
                                      TITLE OF INVENTION: ANTISENSE MODULATION OF HKR1
FILE REFERENCE: RTS-0248
CURRENT APPLICATION NUMBER: US/09/898,556A
CURRENT FILING DATE: 2001-07-03
NUMBER OF SEQ ID NOS: 89
SEQ ID NO 25
LENGTH: 20
TYPE: DNA
ORGANISM: Artificial Sequence
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SEQ ID NO 23
LENGTH: 20
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TITLE OF INVENTION: ANTISENSE MODULATION OF HKR1
FILE REFERENCE: RTS-0248
                                                                                                                                                                                            APPLICANT: C. Frank Bennett APPLICANT: Susan M. Freier
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TITLE OF INVENTION: ANTISENSE MODULATION OF HKR1
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               OTHER INFORMATION: Antisense Oligonucleotide
                              FEATURE:
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ORGANISM: Artificial Sequence
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Pred. No.
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Pred. No.
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1.4e+04;
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TITLE OF INVENTION: ANTISENSE MODULATION OF HK
FILE REFERENCE: RTS-0248
CURRENT APPLICATION NUMBER: US/09/898,556A
CURRENT FILING DATE: 2001-07-03
NUMBER OF SEQ ID NOS: 89
SEQ ID NO 26
LENGTH: 20
TYPE: DNA
ORGANISM: Artificial Sequence
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US-09-898-556A-26/c
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US-09-898-556A-27
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RESULT 39
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                                                                                                                                                                                                                                                                                                                          Publication No. US20030087849A1
GENERAL INFORMATION:
APPLICATION:
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Best Local S
Matches 20
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APPLICANT: C. Frank
APPLICANT: Susan M.
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LENGTH: 20
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Best Local
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                                                                                                                                                                                                                                                                                APPLICANT: C. Frank Bennett
APPLICANT: Susan M. Freier
TITLE OF INVENTION: ANTISENSE MODULATION OF HKR1 EXPRESSION
FILE REFERENCE: RTS-0248
                                                                                                                                                                       TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
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20; Conserv
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100.0%; Pred. No.
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Pred. No.
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                                                                                                                   DB 9;
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US-09-898-556A-28/c

Sequence 28, Application US/09898556A Publication No. US20030087849A1 GENERAL INFORMATION:

APPLICANT: C. Frank Bennett

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Pred. No.

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; OTHER INFORMATION: Antisense Oligonucleotide US-09-898-556A-30
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US-09-898-556A-30/c
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                                                                                                                   SEQ ID NO 30
LENGTH: 20
      Query Match
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                                                                                                                                            FILE REFERENCE: RTS-0248
CURRENT APPLICATION NUMBER: US/09/898,556A
CURRENT FILING DATE: 2001-07-03
NUMBER OF SEQ ID NOS: 89
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
                                                                                                                                                                                                                                APPLICANT: C. Frank Bennett APPLICANT: Susan M. Freier
                                                                                                                                                                                                            TITLE OF INVENTION: ANTISENSE MODULATION OF
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CURRENT FILING DATE: 2001-07-03
NUMBER OF SEQ ID NOS: 89
SEQ ID NO 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQ ID NOS: 89
SEQ ID NO 29
LENGTH: 20
TYPE: DNA
ORGANISM: Artificial Sequence
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TITLE OF INVENTION: ANTISENSE MODULATION OF HKR1 EXPRESSION
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                                                                                ORGANISM: Artificial Sequence
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TITLE OF INVENTION: ANTISENSE MODULATION OF HKR1 EXPRESSION
FILE REFERENCE: RTS-0248
                                                                      FEATURE:
                                                                                                      TYPE: DNA
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TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
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les 20; Conserv
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Pred. No.
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Pred. No.
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Length 20;
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RESULT 44
US-09-898-556A-33/c
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US-09-898-556A-31/c
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GENERAL INFORMATION:
APPLICANT: C. Frank Bennett
APPLICANT: Susan M. Freier
                                          Sequence 33, Application US/09898556A
Publication No. US20030087849A1
                                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                                                            SEQ ID NO 32
LENGTH: 20
TYPE: DNA
ORGANISM: Artificial
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Matches
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Best Local
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LENGTH: 20
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APPLICANT: Susan M. Freier
TITLE OF INVENTION: ANTISENSE MODULATION OF HKR1 EXPRESSION
FILE REFERENCE: RTS-0248
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TITLE OF INVENTION: ANTISENSE MODULATION OF
FILE REFERENCE: RTS-0248
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APPLICANT: Susan M. Freier
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ORGANISM: Artificial
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Pred. No.
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Pred. No.
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RESULT 45
US-09-898-556A-34/c
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US-09-898-556A-35/c
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LENGTH: 20
TYPE: DNA
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Best Local Similarity
Matches 20; Conserv
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APPLICANT: C. Frank
APPLICANT: Susan M
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LENGTH: 20
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SEQ ID NO 35
LENGTH: 20
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Query Match
Best Local Similarity
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                                                                                                                                                                                    FILE REFERENCE: RTS-0248
CURRENT APPLICATION NUMBER: US/09/898,556A
CURRENT FILING DATE: 2001-07-03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: ANTISENSE MODULATION OF HKR1 EXPRESSION FILE REFERENCE: RTS-0248
CURRENT APPLICATION NUMBER: US/09/898,556A
CURRENT FILING DATE: 2001-07-03
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TITLE OF INVENTION: ANTISENSE MODULATION OF HKR1 EXPRESSION
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                                                                                    ORGANISM: Artificial Sequence FEATURE:
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                                                                    OTHER INFORMATION:
                                                                                                                     TYPE: DNA
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Pred. No.
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 Score 20;
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1.4e+04;
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                                                                                                          RESULT 49
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SEQ ID NO 37
LENGTH: 20
TYPE: DNA
ORGANISM: Artificial Sequence
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                                              GENERAL INFORMATION:
                                                           Sequence 38, Application US/09898556A Publication No. US20030087849A1
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Publication No.
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Best Local Similarity 100.0%;
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Best Local
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TITLE OF INVENTION: ANTISENSE MODULATION OF
FILE REFERENCE: RTS-0248
             APPLICANT: C. Frank Bennett
APPLICANT: Susan M. Freier
                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION NUMBER: US/09/898,556A CURRENT FILING DATE: 2001-07-03
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NUMBER OF SEQ ID NOS: 89
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TITLE OF INVENTION: ANTISENSE MODULATION OF
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TITLE OF INVENTION: ANTISENSE MODULATION OF HKR1 EXPRESSION
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TYPE: DNA
ORGANISM: Artificial Sequence
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No. US20030087849A1
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US-09-898-556A-39/c

Sequence 39, Application US/09898556A

Publication No. US20030087849A1

GENERAL INFORMATION:

APPLICANT: C. Frank Bennett

APPLICANT: Susan M. Freiter

TITLE OF INVENTION: ANTISENSE MODULATION OF HKR1 EXPRESSION

FILE REFERENCE: RTS-0248

CURRENT APPLICATION NUMBER: US/09/898,556A

CURRENT APPLICATION NUMBER: US/09/898,556A

CURRENT FILING DATE: 2001-07-03

SEQ ID NO 39

LENGTH: 20

TYDE: DNA

ORGANISM: Artificial Sequence

FEATURE:
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NUMBER OF SEQ ID NOS: 89
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LENGTH: 20
TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Antisense Oligonucleotide
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| 170<br>171<br>171<br>172  | C 167   | c 164   | . 16                                   | 11                                     | c 157                                  |                                | - 11 I                                 | 152<br>153        | c 151                                  | 149               | 147<br>148       | 146                                    | 144               | 142<br>143        | 140<br>141                             | 138                                    |                                      | C 134                                   | 133                                    | 131                                     | 129                                     | c 127             | 126                                      | 123                               | 122                                    |  | C 118                                  |                    |  | ш.                                     |                   | c 110                                      | . د ا              | 106<br>107        | c 104<br>105                             | 101<br>102<br>103  |
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| Sequence 8, Appli<br>Sequence 177, App<br>Sequence 189, App<br>Sequence 177, App                          | 47,   | 6, 26   | 72                                     | 8,                                     |  | 11,                            | 60,                                    | 41,               | 72,<br>14,                             | 72,               | 57,              | 57,<br>72,                             | 57,               | 72,               | 72,                                    | 7,2                                    | 13, App<br>72, App                   | 769,<br>1113,                           | 63                                     | 296,<br>296,                            | 296<br>296                              | 43, A             | 11,                                      | 11, A                             | 11,<br>11,                             | 11, 7                                  | 43,                                    | Sequence 19, April | 1, Ar                                  | Sequence 24, Appl<br>Sequence 24, Appl | 1ce 74            | e 27<br>e 29                               | equence            | equence 5,        | equence 4, App                           | Sequence 194, App<br>Sequence 41, Appl<br>Sequence 13, Appl        |
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| C 243<br>C 243<br>244<br>245  | c 239<br>c 240<br>c 241                                     |   | c 234                                  |  |  | 228<br>229                     |  | c 225             |  | c 222             | 220              | 218<br>219                             |                   |                   |  | c 211<br>c 212                         |                                      |   |  | 203<br>c 204                            |   | c 199<br>c 200    |  | c 196<br>197                      | c 194<br>195                           |  | c 191                                  | c 189              |  | 185<br>186                             | 184               | 182<br>183                                 | 181                | 179               | 177                                      | c 174<br>175<br>176  |
|   | 239 17.<br>240 17.<br>241 17.                               | 236 17.<br>237 17.<br>238 17.                                 | 234 17.<br>235 17.                     | 232 17.<br>233 17.                     | 230 17.<br>231 17.                     | 17<br>17                       | 226 17                                 | 225 17            | 223 17.<br>224 17.                     | 222 17            | 17               | 17                                     | 217 17            | 215 17            | 213 17<br>214 17                       | 211 17<br>212 17                       | 209 17<br>210 17                     | 207 17<br>208 17                        | 205 17<br>206 17                       | 203 17<br>204 17                        | 201 17<br>202 17                        | 200 17            | 198 17                                   | 196 17                            | 194 17<br>195 17                       | 192 17<br>193 17                       | 190 17                                 | 189 . 17           | 187 17                                 | 17<br>17                               | 17                | 17<br>17                                   | 17                 | 17                | 17                                       | 174 17<br>175 17<br>176 17   |
| 242 17.<br>243 17.<br>244 17.<br>245 17.<br>246 17.   | 239 17.6 0<br>240 17.6 0<br>241 17.6 0                      | 236 17.6 0<br>237 17.6 0<br>238 17.6 0                        | 234 17.6 0<br>235 17.6 0               | 232 17.6 0<br>233 17.6 0               | 230 17.6 0<br>231 17.6 0               | 17.6 0<br>17.6 0               | 226 17.6 0                             | 225 17.6 0        | 223 17.8 0<br>224 17.6 0               | 222 17.8 0        | 17.8 0           | 17.8 0<br>17.8 0                       | 217 17.8 0        | 215 17.8 0        | 213 17.8 0<br>214 17.8 0               | 211 17.8 0<br>212 17.8 0               | 209 17.8 0<br>210 17.8 0             | 207 17.8 0<br>208 17.8 0                | 205 17.8 0<br>206 17.8 0               | 203 17.8 0<br>204 17.8 0                | 201 17.8 0<br>202 17.8 0                | 200 17.8 0        | 198 17.8 0                               | 196 17.8 0                        | 194 17.8 0<br>195 17.8 0               | 192 17.8 0<br>193 17.8 0               | 190 17.8 0                             | 189 . 17.8 0       | 187 17.8 0                             | 17.8 C                                 | 17.8              | 17.8 C                                     | 17.8               | 17.8              | 17.8                                     | 174 17.8 (<br>175 17.8 (<br>176 17.8 (                             |
| 242 17.6 0.6 43<br>243 17.6 0.6 43<br>244 17.6 0.6 45<br>245 17.6 0.6 45<br>246 17.6 0.6 45               | 239 17.6 0.6 39<br>240 17.6 0.6 41<br>241 17.6 0.6 42       | 23b 17.6 0.6 36<br>237 17.6 0.6 36<br>238 · 17.6 0.6 37       | 234 17.6 0.6 36<br>235 17.6 0.6 36     | 232 17.6 0.6 36<br>233 17.6 0.6 36     | 230 17.6 0.6 36<br>231 17.6 0.6 36     | 17.6 0.6 35<br>17.6 0.6 35     | 226 17.6 0.6 35<br>227 17.6 0.6 35     | 225 17.6 0.6 28   | 223 17.8 0.6 60<br>224 17.6 0.6 27     | 222 17.8 0.6 59   | 17.8 0.6 58      | 17.8 0.6 57<br>17.8 0.6 58             | 217 17.8 0.6 57   | 215 17.8 0.6 54   | 213 17.8 0.6 54<br>214 17.8 0.6 54     | 211 17.8 0.6 54<br>212 17.8 0.6 54     | 209 17.8 0.6 54<br>210 17.8 0.6 54   | 207 17.8 0.6 54<br>208 17.8 0.6 54      | 205 17.8 0.6 53<br>206 17.8 0.6 54     | 203 17.8 0.6 53<br>204 17.8 0.6 53      | 201 17.8 0.6 52<br>202 17.8 0.6 53      | 200 17.8 0.6 50   | 198 17.8 0.6 49                          | 196 17.8 0.6 49                   | 194 17.8 0.6 49<br>195 17.8 0.6 49     | 192 17.8 0.6 47<br>193 17.8 0.6 47     | 190 17.8 0.6 47<br>191 17.8 0.6 47     | 189 17.8 0.6 47    | 187 17.8 0.6 47                        | 17.8 0.6 47<br>17.8 0.6 47             | 17.8 0.6 45       | 17.8 0.6 45                                | 17.8 0.6 45        | 17.8 0.6 45       | 17.8 0.6 43                              | 174 17.8 0.6 38<br>175 17.8 0.6 38<br>176 17.8 0.6 43              |
| 242 17.6 0.6<br>243 17.6 0.6<br>244 17.6 0.6<br>245 17.6 0.6 .  | 239 17.6 0.6 39 4<br>240 17.6 0.6 41 1<br>241 17.6 0.6 42 4 | 23b 17.6 0.6 36 3<br>237 17.6 0.6 36 4<br>238 · 17.6 0.6 37 4 | 234 17.6 0.6 36 2<br>235 17.6 0.6 36 2 | 232 17.6 0.6 36 2<br>233 17.6 0.6 36 2 | 230 17.6 0.6 36 1<br>231 17.6 0.6 36 1 | 17.6 0.6 35 4<br>17.6 0.6 35 4 | 226 1/.6 0.6 35 1<br>227 17.6 0.6 35 4 | 225 17.6 0.6 28 1 | 223 17.8 0.6 60 3<br>224 17.6 0.6 27 4 | 222 17.8 0.6 59 2 | 17.8 0.6 58 4    | 17.8 0.6 57 4<br>17.8 0.6 58 1         | 217 17.8 0.6 57 1 | 215 17.8 0.6 54 4 | 213 17.8 0.6 54 3<br>214 17.8 0.6 54 4 | 211 17.8 0.6 54 2<br>212 17.8 0.6 54 3 | 209 17.8 0.6 54 1<br>210 17.8 0.6 54 | 207 17.8 0.6 54 1<br>208 17.8 0.6 54 1  | 205 17.8 0.6 53 4<br>206 17.8 0.6 54 1 | 203 17.8 0.6 53 4<br>204 17.8 0.6 53 4  | 201 17.8 0.6 52 4<br>202 17.8 0.6 53 4  | 200 17.8 0.6 51 4 | 198 17.8 0.6 49 4                        | 196 17.8 0.6 49 4                 | 194 17.8 0.6 49 4<br>195 17.8 0.6 49 4 | 192 17.8 0.6 47 4<br>193 17.8 0.6 47 4 | 190 17.8 0.6 47 2<br>191 17.8 0.6 47 3 | 189 17.8 0.6 47 1  | 187 17.8 0.6 47 1                      | 17.8 0.6 47 1<br>17.8 0.6 47 1         | 17.8 0.6 45 4     | 17.8 0.6 45 4                              | 17.8 0.6 45 4      | 17.8 0.6 45 4     | 17.8 0.6 43 1                            | 174 17.8 0.6 38 1<br>175 17.8 0.6 38 4<br>176 17 8 0.6 43 1        |

| C 315<br>C 317<br>C 317<br>C 317<br>C 317<br>C 317   | 308<br>309<br>c 310<br>311  | 304<br>305<br>307   | 302   | 29 <b>8</b><br>299  |  | c 294<br>c 295                          |  |                                    |  | C 286   |  | 280<br>281<br>282   | c 278<br>279   | 276                                    |   | c 271<br>c 272                         |  |  | c 265<br>266                           |  | 261<br>c 262                            | c 259<br>260                           | SS                                     | ·σι                                    | ח טונ                          | וסות                                   | л UI -                              | 247<br>248<br>249                       |
|--|---|---|---|---|--|---|--|------------------------------------|--|---|--|---|--|--|---|--|--|--|--|--|---|--|--|--|--------------------------------|--|-------------------------------------|---|
| 17.4<br>17.4<br>17.4<br>17.4<br>17.4   |   | 17.4  | 17.4  | 17.4<br>17.4  | 17.4                                   | 17.4                                    | 17.4                                   | 17.4                               | 17.4                                       | 17.4  | 17.6   | 17.6<br>17.6  |  |  | 17.6  |  | 17.6<br>17.6                           |  |  |  |   | 17.6<br>17.6                           |  | 17.6                                   |                                |  | · · ·                               | 17.6<br>17.6                            |
| 00000000   |   |   |   |   |  |   |  |                                    |  |   |  |   |  |  |   |  |  |  |  |  |   |  |  |  |                                |  |                                     |   |
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| US-08-495-6958-20<br>US-08-343-682-3<br>US-08-343-682-6<br>US-08-652-558-28<br>US-08-171-389-4<br>US-08-173-936-4<br>US-08-475-228A-4                | US-09-290-338-9<br>US-09-464-453-131<br>US-09-117-860-44<br>US-09-117-860-46<br>US-09-117-860-46      | US-09-290-577-9<br>US-09-290-452-8<br>US-09-290-452-9<br>US-09-290-452-9<br>US-09-290-338-8 | US-08-986-617-12<br>US-08-813-507-131<br>US-09-290-577-8  | US-07-981-525-12<br>US-08-220-033-12<br>US-08-465-491-12                    | US-08-435-628-181<br>US-08-962-281-18  | US-08-702-870A-17<br>US-08-373-124A-181 | US-09-038-073-1559 US-08-273-362-1     | US-09-071-845-1408                 | US-08-292-620A-1505<br>US-08-585-684B-1559 | US-08-280-228-6<br>US-08-291-932A-516<br>US-08-293-630A-1408  | PCT-US96-03916-49<br>US-07-618-312A-6  | US-08-754-681-19<br>US-09-357-487B-19<br>US-09-602-428-19 | US-08-431-32/A-/<br>US-08-053-451B-121<br>US-08-911-894-64 | PCT-US92-10024-8                       | US-09-537-911A-26<br>US-09-710-200-13<br>US-09-423-430-13 | US-08-290-793B-52<br>US-08-284-516C-26 | US-08-704-744-52<br>US-08-469-557-52   | PCT-US96-06053-10<br>US-08-208-886C-52 | US-08-906-156A-71<br>PCT-US95-13975-59 | US-08-646-695-10<br>US-09-171-025-21   | US-08-207-901-78<br>US-08-860-174A-22   | US-08-849-567A-69<br>US-09-206-942-95  | US-09-641-638-1167<br>US-08-756-506-18 | US-08-222-177A-458                     | PCT-US93-12388-185             | US-09-069-821-23.                      | US-08-864-473-70                    | US-08-475-228A-185<br>US-08-452-164A-37 |
| equence 2 equence 3 equence 6 equence 6 equence 2 equence 4 equence 4  |   | ာတထာတဏ  | ייים  | 12,   | 181, App                               | ب بر ب                                  | 155                                    | 14(                                | 150  | د ال م  | 6,4  | Sequence 19, Appl<br>Sequence 19, Appl                    | 121,   | 7 × ×                                  | 4 14 6  | 26,                                    | 52,                                    | 10,<br>52,                             | 71,                                    | 10,                                    | 78,                                     | 10 M                                   | 116                                    | 458                                    | 185                            | 23 2                                   | uence 70,                           | ce 185,<br>ce 37,                       |
|  |   |   |   |   |  |   |  |                                    |  |   |  |   |  |  |   |  |  |  |  |  |   |  |  |  |                                |  |                                     |   |
|  |   | -   | •   | •   | -                                      |   |  |                                    |  |   |  |   |  |  |   |  |  |  |  |  |   |  |  |  |                                |  |                                     |   |
|  | n   |   | a (   |   |  | o 0                                     | 0.0                                    |                                    |  | ) n   | 0  | c   |  | a                                      |   | C                                      |  | o o                                    |  | 200                                    | <b>.</b>                                | c                                      | 0                                      | Ω                                      |                                | o c                                    | ) O (                               | n 0                                     |
| C C 388<br>C C 388<br>389<br>390<br>391  |   |   |   |   | to to t                                |   | ( (.                                   |                                    | (.) (.                                     |   | (a) (a) (  | a ta ta   | 350<br>351<br>352  | 3 (4) (4                               |   | ta ta                                  |  |  |  |  |   |  |  |  | 326                            |  |                                     | c 320<br>321                            |
| 3 3 8 8 8 7 1 1 3 3 8 8 9 1 1 1 3 3 9 2 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1  | 381 17.<br>382 17.<br>383 17.<br>384 17.  | 377 17.<br>378 17.<br>379 17.<br>379 17.  | 374 17.<br>375 17.  | 371 17.<br>372 17.  | 369 17:<br>370 17.                     | 367 17.                                 | 365 17                                 | 363 17.                            | 361 17.                                    | 358 17.<br>359 17.  | 356 17.<br>357 17.   | 353 17.<br>354 17.  | 50 17.<br>51 17.<br>52 17.                                 | 348 1/.<br>349 17.                     | 346 17.<br>347 17.  | 344 17.<br>345 17.                     | 342 17.<br>343 17.                     | 340 17.<br>341 17.                     | 338 17.<br>339 17.                     | 336 17.                                | 334 17.                                 | 332 17.<br>333 17.                     | 330 17.<br>331 17.                     | 328 17.                                | 17.                            | 324 17.<br>325 17.                     | 323 17.                             | 320 17.<br>321 17.                      |
| 385 17. 386 17. 387 17. 388 17. 389 17. 390 17. 391 17. 392 17.  | 381 17.2<br>382 17.2<br>383 17.2  | 377 17.2<br>378 17.2<br>379 17.2  | 374 17.2 (<br>375 17.2 (  | 371 17.2 (<br>372 17.2 (<br>373 17.2 (                                      | 369 17:2 (<br>370 17.2 (               | 367 17.2 C                              | 365 17.2                               | 363 17.2                           | 361 17.2 (<br>361 17.2 (                   | 358 17.2 (<br>359 17.2 (  | 356 17.2 (357 17 | 353 17.2 (<br>354 17.2 (                                  | 50 17.2 (<br>51 17.2 (<br>52 17.2 (                        | 349 17.2                               | 346 17.4 (<br>347 17.4 (                                  | 344 17.4 C                             | 342 17.4 C                             | 340 17.4 C                             | 338 17.4 C                             | 336 17.4                               | 334 17.4                                | 332 17.4 C                             | 330 17.4 C                             | 328 17.4 (                             | 17.4                           | 324 17.4 (                             | 323 17.4 (                          | 320 17.4 C                              |
| 385 17.2 0.6 44<br>387 17.2 0.6 44<br>387 17.2 0.6 45<br>389 17.2 0.6 45<br>389 17.2 0.6 45<br>390 17.2 0.6 46<br>391 17.2 0.6 46<br>392 17.2 0.6 46 | 381 17.2 0.6 39<br>382 17.2 0.6 44<br>383 17.2 0.6 44<br>384 17.2 0.6 44                              | 377 17.2 0.6 38<br>378 17.2 0.6 38<br>379 17.2 0.6 38<br>379 17.2 0.6 38                    | 374 17.2 0.6 38<br>375 17.2 0.6 38<br>376 17.2 0.6 38   | 371 17.2 0.6 36<br>372 17.2 0.6 38<br>373 17 2 0.6 38                       | 369 17:2 0.6 36<br>370 17.2 0.6 36     | 367 17.2 0.6 34<br>368 17.2 0.6 36      | 365 17.2 0.6 34<br>366 17.2 0.6 34     | 363 17.2 0.6 34<br>364 17.2 0.6 34 | 360 17.2 0.6 33<br>361 17.2 0.6 33         | 358 17.2 0.6 30<br>359 17.2 0.6 33  | 356 17.2 0.6 30<br>357 17.2 0.6 30   | 353 17.2 0.6 28<br>354 17.2 0.6 30                        | 50 17.2 0.6 25<br>51 17.2 0.6 25<br>52 17.2 0.6 28         | 348 17.4 0.6 60<br>349 17.2 0.6 24     | 346 17.4 0.6 60<br>347 17.4 0.6 60                        | 344 17.4 0.6 59<br>345 17.4 0.6 60     | 342 17.4 0.6 58<br>343 17.4 0.6 58     | 340 17.4 0.6 57<br>341 17.4 0.6 58     | 338 17.4 0.6 57<br>339 17.4 0.6 57     | 336 17.4 0.6 57                        | 334 17.4 0.6 53<br>335 17.4 0.6 54      | 332 17.4 0.6 52<br>333 17.4 0.6 53     | 330 17.4 0.6 52<br>331 17.4 0.6 52     | 328 17.4 0.6 52<br>329 17.4 0.6 52     | 17.4 0.6 49                    | 324 17.4 0.6 47<br>325 17.4 0.6 47     | 323 17.4 0.6 47                     | 320 17.4 0.6 47<br>321 17.4 0.6 47      |
| 386 17.2 0.6 4386 17.2 0.6 4388 17.2 0.6 4389 17.2 0.6 4390 17.2 0.6 4391 17.2 0.6 4391 17.2 0.6 4391 17.2 0.6 44                                    | 381 17.2 0.6 39 4<br>382 17.2 0.6 44 1<br>383 17.2 0.6 44 1<br>384 17.2 0.6 44 1<br>386 17.2 0.6 44 1 | 377 17.2 0.6 38 1<br>378 17.2 0.6 38 5<br>378 17.2 0.6 38 5<br>380 17.2 0.6 38 5            | 374 17.2 0.6 38 1 US-08-599-252-74 37 38 1 US-08-599-252-74 38 1 US-08-436-074-47 375 17.2 0.6 38 1 US-08-436-074-47 | 371 17.2 0.6 36 4 US-08-505-860C-12<br>372 17.2 0.6 38 1 US-08-373-124A-138 | 369 17:2 0.6 36 2<br>370 17.2 0.6 36 3 | 367 17.2 0.6 34 4<br>368 17 2 0.6 36 2  | 365 17.2 0.6 34 4<br>366 17.2 0.6 34 4 | 363 17.2 0.6 34 3                  | 360 17.2 0.6 33 5<br>361 17.2 0.6 33 5     | 358 17.2 0.6 30 4 US-08-642-274D-19 358 17.2 0.6 33 1 US-08-292-945-1 359 17.2 0.6 33 1 US-08-25-23-1 | 356 17.2 0.6 30 2<br>357 17.2 0.6 30 2   | 353 17.2 0.6 28 4<br>354 17.2 0.6 30 1                    | 50 17.2 0.6 25 4<br>51 17.2 0.6 25 4<br>52 17.2 0.6 28 1   | 349 17.2 0.6 24 4<br>349 17.2 0.6 24 4 | 346 17.4 0.6 60 3<br>347 17.4 0.6 60 4                    | 344 17.4 0.6 59 4<br>345 17.4 0.6 60 3 | 342 17.4 0.6 58 2<br>343 17.4 0.6 58 3 | 340 17.4 0.6 57 4<br>341 17.4 0.6 58 1 | 338 17.4 0.6 57 3<br>339 17.4 0.6 57 4 | 336 17.4 0.6 57 1<br>337 17.4 0.6 57 1 | 334 17.4 0.6 53 1<br>335 17.4 0.6 54 2. | 332 17.4 0.6 52 2<br>333 17.4 0.6 53 1 | 330 17.4 0.6 52 2<br>331 17.4 0.6 52 2 | 328 17.4 0.6 52 I<br>329 17.4 0.6 52 1 | 17.4 0.6 49 1<br>17.4 0.6 49 5 | 324 1/.4 0.6 4/ 4<br>325 17.4 0.6 47 5 | 323 17.4 0.6 47 4 US-09-641-638-820 | 320 17.4 0.6 47 3<br>321 17.4 0.6 47 4  |

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|--|---|---|---|---|--|--|---|--|--|---|--|---|--|--|---|---|--|---|---|--|---|---|---|--|
| 17717171717  | 17.2<br>17<br>17<br>17<br>17  | 17.2<br>17.2<br>17.2  |   | 17.2<br>17.2<br>17.2                                |  |  |   | 17.2<br>17.2<br>17.2                                       |  | 17.2<br>17.2  |  | 17.2  | 17.2   | 17.2<br>17.2                               |   |   |  |   | 17.2<br>17.2                                | 17.2<br>17.2                                 | 17.2<br>17.2                              | 17.2<br>17.2                                | 17.2<br>17.2  | 17.2<br>17.2   |
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| & & & & & & & & & & & & & & & & & & &  | 100000  | 50000   | 000   | O W W   | 000  | 154                                    | ωω Ν ι  | 2000   | 000  | 000   | 000  | 0 40  | o wo a   | 000  | <b>&amp;</b> &                            | ထထ                                      | ∞ σο ¬   | 7 7   | 7   | 7  |   |   |   |  |
|  |   | ,444,   |   | 200   | .4.4   | . 4 4 .                                |   | 1000   | 440  | 321   |  | -4  | 3 H G  |  | 22  | 22                                      |  | л 4- г                                      | 4 4   | ω <b>4</b>                                   | ω N                                       | 21  |   | <br>   |
| US-08-311-486C-797<br>US-08-311-486C-798<br>US-08-585-684B-2021<br>US-08-585-684B-2022<br>US-09-038-073-2021<br>US-09-038-073-2022<br>US-09-737-698B-16<br>US-08-106-761-9<br>US-08-327-874A-33<br>PCT-1FS4A-09700-33  | 15-09-218-089A-11<br>15-09-218-089A-11<br>15-08-584-040-3686<br>15-09-033-055A-17<br>15-08-106-761-5<br>15-08-451-715A-53                               | JS-09-055-075C-7<br>JS-09-055-075C-9<br>JS-09-055-075C-11<br>JS-09-055-075C-115 | JS-08-11/-932-760<br>JS-08-11/-952-760<br>JS-07-921-104D-25 | 5422249-10<br>JS-08-704-473-21<br>JS-08-117-952-759 | JS-08-129-/22A-/<br>JS-08-219-012-55<br>JS-08-687-421-243                | US-08-797-812-5<br>US-09-357-740-18    | JS-00-9/3-310-19<br>JS-09-234-874A-68<br>JS-08-429-181-40<br>JS-08-164-388-40 | US-08-222-177A-337<br>US-08-829-876-68<br>US-08-943-136-19 | JS-08-687-421-5<br>JS-09-354-947-558<br>PCT-US93-12388-558                                 | JS-08-123-936-558<br>JS-08-475-228A-558<br>JS-08-482-080A-558   | US-08-171-389-558<br>US-08-384-708A-5  | US-09-234-874A-17<br>US-07-828-444-6  | US-08-475-680-67<br>US-08-198-670A-34<br>US-08-838-876-17                                | US-08-714-017-67<br>US-07-602-848E-13      | US-08-483-932-67<br>US-08-720-420A-67     | US-08-487-113D-67<br>US-08-473-503-67   | US-08-482-882-67<br>US-08-483-389-67   | US-09-641-638-974                           | US-09-641-638-952                           | US-08-482-080A-237<br>US-09-354-947-237      | US-08-771-624B-16<br>US-08-155-938-2      | US-08-123-936-237<br>US-08-475-228A-237     | US-09-249-215-113<br>US-08-171-389-237  | US-08-910-408-113<br>US-08-803-085-19<br>US-09-249-215-90                                  |
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| ) W > 0000000  | 1, Appl<br>686, Ap<br>7, Appl<br>7, Appl<br>3, Appl   |   | 11000   | 2 11 12   |  | æ · c                                  | 2004  | ωφωι   | 558,<br>558,   | ن ن ن   | · 0:   | 7   | 147  | 37   | 77.                                       | 77,                                     | 770  | 2 -1 0                                      | יוט ע                                       | - بن بن                                      | 0   |   | $\omega \vdash c$   | · ω  |
| * <del>2</del>   | ים<br>יסא סיוטי   | P P P P   | App<br>App<br>Pp1   | 249<br>pp1  | pp1  | 100                                    | 10.0  | 7, Appl  | pili<br>App  | Alp<br>Alp<br>Alp   | App. i   | 19.0  | 125  | P P  | 1,10                                      |   |  | App   | App   | App  | ppl<br>lgg                                | App   | App   | App<br>pp1   |
| App App App App 11 App App 11 App App 11 App 12 App | יים פים ביק   |   |   |   |  |  |   |  |  |   |  |   |  |  |   |   |  |   |   |  |   |   |   |  |
| PETER PROPERTY OF THE PROPERTY | COMPE   | STA COMPU   | ; ADD; ; STH  | NUMBE   | ; APPLIC<br>; TITLE<br>; TITLE   | ; Sequence<br>; Patent No<br>; GENERAL | ULT 1<br>08-294   |  | c 499<br>c 500   | 496<br>C 497<br>C 498   | 494<br>494<br>495  |   | 489<br>490   |  | C 485                                     |   | . c 481  |   | c 477                                       |  | c 473                                     | 471   | 468<br>469<br>6 470   | າຫ້ອ   |
| PETER PROPERTY OF THE PROPERTY | COMPE   | STA COMPU   | ; ADDRESSE<br>; STREET:<br>; CITY: B                        | ; NUMBER  |  | tent No                                | SULT 1<br>-08-294-4   |  | 499  | ~ ~ ~   | 494<br>494<br>495  | 492   | 90   | 487  | 485                                       | 483                                     | 481  | 479   | 477   | 475  |   | » н (                                       | 70<br>70  | 466  |
| pp ; FILING DATE: CLASSIFICATION CLASSIFICATION APPLICATION APPLIC | COMPE   | STA COMPU   | ; ADDRESSEE:<br>; STREET: 22<br>; CITY: Bost                | ; NUMBER OF SEQ<br>; CORRESPONDENC                  |  | tent No                                | SULT 1<br>-08-294-4   |  | 499 17 0.<br>500 17 0.   | 496<br>497<br>498   | 494 17 0.<br>495 17 0.   | 491 17 0.<br>492 17 0.  | 89 17 0.<br>90 17 0.   | 487 17 0.<br>488 17 0.                     | 485 17 0.<br>486 17 0.                    | 483 17 0                                | 480 17 0.<br>481 17 0.   | 479 17 0.                                   | 477 17 0                                    | 475 17 0                                     | 17 0                                      | 1 17 0                                      | 70 17 0   | 466 17 0<br>467 17 0   |
| pp ; FILING DATE: CLASSIFICATION Ap ; PRIOR APPLICATION NU ANME: Freeman   | COMPOUTER: 3.5 UT  COMPOUTER: IBM PS/2 M  OPERATING SYSTEM: IB  SOFTWARE: WordPerfec  CURRENT APPLICATION NUMBER:                                       | STATE: Massachuse COUNTRY: U.S.A. ZIP: 02110-2804 COMPUTER READABLE FO          | ; ADDRESSEE:<br>; STREET: 22<br>; CITY: Bost                | NUMBER OF SEQUENCES: CORRESPONDENCE ADDRESS         | CANT: Vary, Cal<br>OF INVENTION:<br>OF INVENTION:                        | tent No                                | SULT 1<br>-08-294-4   |  | 499 17 0.6 5<br>500 17 0.6 5   | 496 17 0.<br>497 17 0.<br>498 17 0.   | 494 17 0.6 4<br>495 17 0.6 4   | 491 17 0.6 4<br>492 17 0.6 4  | 89 17 0.6 4<br>90 17 0.6 4   | 487 17 0.6 4<br>488 17 0.6 4               | 485 17 0.6 4<br>486 17 0.6                | 483 17 0.6                              | 480 17 0.6 4<br>481 17 0.6 4   | 479 17 0.6                                  | 477 17 0.6                                  | 475 17 0.6                                   | 17 0.6                                    | 1 17 0.6                                    | 58 17 0.6 4<br>69 17 0.6 4  | 466 17 0.6 4<br>467 17 0.6   |
| pp ; FILING DATE: CLASSIFICATION CLASSIFICATION APPLICATION NO APPLICATION NO APPLICATION  | COMPOUTER: 3.5 UT  COMPOUTER: IBM PS/2 M  OPERATING SYSTEM: IB  SOFTWARE: WordPerfec  CURRENT APPLICATION NUMBER:                                       | STATE: Massachuse COUNTRY: U.S.A. ZIP: 02110-2804 COMPUTER READABLE FO          | ; ADDRESSEE:<br>; STREET: 22<br>; CITY: Bost                | NUMBER OF SEQUENCES: CORRESPONDENCE ADDRESS         | CANT: Vary, Cal<br>OF INVENTION:<br>OF INVENTION:                        | tent No                                | SULT 1<br>-08-294-4   |  | 499 17 0.6 51 4<br>500 17 0.6 52 3   | 496: 17 0.6 49 4<br>497 17 0.6 50 4<br>498 17 0.6 51 2  | 493 17 0.6 48 4<br>494 17 0.6 49 2<br>495 17 0.6 49 4  | 491 17 0.6 48 2<br>492 17 0.6 48 4<br>493 17 0.6 48 4                               | 89 17 0.6 47 4<br>90 17 0.6 47 4   | 487 17 0.6 47 4<br>488 17 0.6 47 4         | 485 17 0.6 45 4<br>486 17 0.6 46 4        | 483 17 0.6 45 4<br>484 17 0.6 45 4      | 480 17 0.6 45 4<br>481 17 0.6 45 4<br>482 17 0.6 45 4  | 479 17 0.6 45 2                             | 477 17 0.6 45 2                             | 475 17 0.6 45 1                              | 17 0.6 44 4                               | 1 17 0.6 44 2                               | 0.6 43 4<br>69 17 0.6 43 4<br>70 17 0.6 44 2  | 466 17 0.6 42 3<br>467 17 0.6 43 4   |
| PP FILING DATE: CLASSIFICATION: 435 PRIOR APPLICATION DATA: APPLICATION NUMBER: APPLICATION NUMBER: APPLICATION NUMBER: FILING DATE: 16 JAN APPLICATION NUMBER: FILING DATE: 17-DEC- PILING DATE: 17-DEC- PILING DATE: FIGURE TORNEY. AFTORNEY.AGENT INFORMAT PILING DATE: FIGURE TORNEY. AMME: FIGURE TORNEY.   | COMPUTER: IBM PS, OPERATING SYSTEM: SOFTWARE: WORDED: CURRENT APPLICATION NUMBER  | STATE: Massachusetts COUNTRY: U.S.A. ZIP: 02110-2804 COMPUTER READABLE FORM:    | ; ADDRESSE<br>; STREET:<br>; CITY: B                        | NUMBER OF SEQUENCES: 49 CORRESPONDENCE ADDRESS:     | CANT: Vary, Calvin OF INVENTION: NUCLEIC ACID OF INVENTION: TRIPLE HELIX | tent No                                | SULT 1<br>-08-294-4   | ALIGNMENTS   | 499 17 0.6 51 4<br>500 17 0.6 52 3   | 496: 17 0.6 49<br>497 17 0.6 50<br>498 17 0.6 51  | 493 17 0.6 48 4<br>494 17 0.6 49 2<br>495 17 0.6 49 4  | 491 17 0.6 48 2<br>492 17 0.6 48 4<br>493 17 0.6 48 4                               | 89 17 0.6 47 4<br>90 17 0.6 47 4   | 487 17 0.6 47 4<br>488 17 0.6 47 4         | 485 17 0.6 45 4<br>486 17 0.6 46 4        | 483 17 0.6 45 4<br>484 17 0.6 45 4      | 480 17 0.6 45 4<br>481 17 0.6 45 4<br>482 17 0.6 45 4  | 479 17 0.6 45 2                             | 477 17 0.6 45 2                             | 475 17 0.6 45 1                              | 17 0.6 44 4                               | 1 17 0.6 44 2                               | 0.6 43 4<br>69 17 0.6 43 4<br>70 17 0.6 44 2  | 466 17 0.6 42 3<br>467 17 0.6 43 4   |
| PP FILING DATE: CLASSIFICATION PRIOR APPLICATION NU | COMPUTER: IBM PS/2 Model 502 or OPERATING SYSTEM: IBM P.C. DOS SOFTWARE: WordPerfect (Version CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/294,4 | STATE: Massachusetts COUNTRY: U.S.A. ZIP: 02110-2804 COMPUTER READABLE FORM:    | ; ADDRESSEE:<br>; STREET: 22<br>; CITY: Bost                | NUMBER OF SEQUENCES: 49 CORRESPONDENCE ADDRESS:     | CANT: Vary, Cal<br>OF INVENTION:<br>OF INVENTION:                        | tent No                                | SULT 1<br>-08-294-4   | ALIGNMENTS   | 499 17 0.6 51 4 US-09-085-028-7 Sequence 7, 500 17 0.6 52 3 US-08-388-029A-31 Sequence 31. | 496: 17 0.6 49 4 US-09-538-709-1004 Sequence<br>497 17 0.6 50 4 US-08-974-911-14 Sequence<br>498 17 0.6 51 2 US-08-525-506-7 Sequence | 494 17 0.6 49 2 US-09-018-914-23<br>494 17 0.6 49 2 US-09-002-177-18<br>495 17 0.6 49 4 US-09-374-584-18 | 491 17 0.6 48 2 US-08-618-911-10 Sequence<br>492 17 0.6 48 US-09-710-200-7 Sequence | 89 17 0.6 47 4 US-09-641-638-1047 Sequence<br>90 17 0.6 47 4 US-09-641-638-1080 Sequence | 487 17 0.6 47 4 US-08-976-183A-10 Sequence | 485 17 0.6 45 4 US-08-977-378-19 Sequence | 483 17 0.6 45 4 US-09-301-03-2 Sequence | 480 17 0.6 45 4 US-08-887-3538-12 Sequence 17<br>481 17 0.6 45 4 US-09-301-153-1 Sequence 17<br>482 17 0.6 45 4 US-00-301-153-2 Sequence 2 | 479 17 0.6 45 2 US-08-882-756-5 Sequence 5, | 477 17 0.6 45 2 US-08-882-756-1 Sequence 1, | 475 17 0.6 45 1 US-08-337-278-60 Sequence 58 | 17 0.6 44 4 US-09-192-6578-27 Sequence 27 | 1 17 0.6 44 2 US-08-790-963-86 Sequence 86. | 58 17 0.5 43 4 US-09-042-353-401 Sequence 401.<br>69 17 0.6 43 4 US-08-758-417A-251 Sequence 251.<br>70 17 0.6 44 2 US-08-331-380A-27 Sequence 27 | 466 17 0.6 42 3 US-08-997-758-4 Sequence 4<br>467 17 0.6 43 4 US-08-960-190A-13 Sequence 1 |

REFERENCE/DOCKET NUMBER:

00088-037001

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US-08-388-029A-30/c
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Best Local Similarity
                                            Matches
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TELEX: 200154
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                            TELEX: AMERPAT INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                           MOLECULE TYPE: DN
                                                                                                              MOLECULE HYPOTHETICAL: P
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                                                                                                                                                                                                                                                                       TELECOMMUNICATION INFORMATION: TELEPHONE: 703-684-1111
                                                                                                                                                                                                                                                                                                                                         ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQUENCES: 9
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: GRANSTROM, DAVID R.
APPLICANT: GAJADHAR, ALVIN A.
TITLE OF INVENTION: SARCOCYSTIS NEURONA DIAGNOSTIC PRIMER
                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: FILING DATE: 14-FEE
                                          Local Similarity 58.7 nes 27; Conservative
                                                                                                                                                                   STRANDEDNESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY:
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                                                                                                                                                          TOPOLOGY:
                                                                                                                                                                                   TYPE: nucleic acid
                                                                                                                                                                                                LENGTH:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          MEDIUM TYPE:
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5. 6110665
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Pred. No. 1
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GENERAL I
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                                                          Matches
                                                                                     Query Match
                                                                          Best Local Similarity
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                                                                                                                                                  ANTI-SENSE: NO ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: GRAY, IAN C
APPLICANT: STEWART, LORNA M
                                                                                                                                                                                                                                                                                                       TELECOMMUNICATION INFORMATION: TELEPHONE: 703-816-4000
                                                                                                                                                                                                                                                                                                                                                                     FILING DATE: 22-OCT-1996
ATTORNEY/AGENT INFORMATION:
NAME: SADOFF, B.J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION DATA: APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS
                                                                                                                                                                             HYPOTHETICAL:
                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE: 23-OCT-1995 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CITY: ARLINGTON STATE: VA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US 60/042,655 FILING DATE: 02-APR-1996
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                                                                                                                                                                                                                                                                                                                                                      REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER:
                                                                                                                                                                                                STRANDEDNESS:
TOPOLOGY: li
                                                                                                                                                                                                                                                                                                                                    REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE:
                                                                                                                                       ORGANISM:
                                                                                                                                                                                                                                                                                                      TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER:
                                                                                                                                                                                                                                             LENGTH:
                              528 CTGTTTTCAAGTTTATGGGCAGGAAATCC 556
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   50 CCGGAANCGAACCCTAATNCCCCNNNACCNNNCACNGCCACGGGAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          22201
                                                                                                                                                                                                                           nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Application US/08906156A
                                                                                                                                                                                                                                             52 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1100 NORTH GLEBE ROAD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        USA
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                                                                                                                                                                                                                                                                                         703-816-4100
                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         E: Floppy disk
IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                linear
                                                                                                                                                                                                                                                                                                                                                                                                    22-OCT-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           JMBER: US 60/033,147
13-DEC-1996
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                                                                         0.8%;
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                                                                                                                                                                                                                                                                                                                                                                                                                  WO PCT/96GB/02588
                                                                                                                                                                                                                                                                                                                                                  36,663
                                                                                                                                                                                                                                                                                                                                      1090-14
                                                          0;
                                                                         Score 21;
Pred. No.
                                                          Mismatches
                                                                       2.6e+03;
13
                                                                                         DB 4;
                                                                                       Length 52;
                                                          Indels
                                                          0;
                                                          Gaps
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RESULT 4 US-08-952-793-236

0;

Sequence 236, Application US/08952793

Patent No. 6280932

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RESULT 5
PCT-US96-09455A-236
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                                                                                                                                                                                                                                                                    ; FEATURE:
; OTHER INFORMATION:
US-08-952-793-236
Sequence 236, Application PC/TUS9609455A
GENERAL INFORMATION:
APPLICANT: PARMA, et al.
TITLE OF INVENTION: HIGH AFFINITY NUCLEIC ACID
TITLE OF INVENTION: LIGANDS TO LECTINS
                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                      Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                TELEFAX: (303) 793-343
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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TITLE OF INVENTION: TO I
NUMBER OF SEQUENCES: 39(
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                             MOLECULE TYPE: RNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            REFERENCE/DOCKET NUMBER: NETELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: 08/47:
FILING DATE: 07-JUNE-1995
ATTORNEY/AGENT INFORMATION:
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APPLICATION NUMBER: 08/47:
PILING DATE: 07-JUNE-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE: 07-JUNE-1995 PRIOR APPLICATION DATA:
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MEDIUM TYPE: Diskette, 3 1/2 diskette, 1.44 MB
                                                                                                                                                                                                                                                                                                             OTHER INFORMATION:
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TOPOLOGY: linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: Barry J. Swanson REGISTRATION NUMBER: 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: FILING DATE: 07-JU
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: PCT/US96/09455
FILING DATE: 05-JUNE-1995
                                                                                                                                                                                                                                                                                                                                                                                                       ENGTH:
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STATE: Colorado
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE:
                                                                                                                                                                140 CTTCTCCAGGAAGAGCACTCAGGAGACCAGGAAAATGGCTACAGG 184
                                                                                                                                  nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                    59 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               E: Swanson & Bratschun, L.L.C. 8400 E. Prentice Avenue, Suite
                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PARMA, et al
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (303)
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                                                                                                                                                                                                                                                                                                                                                                       single
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S: 390
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                                                                                                                                                                                                                 Score 21; DB 4;
Pred. No. 2.9e+03;
                                                                                                                                                                                                  Mismatches
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                                                                                                                                                         Sequence 7, Applic Patent No. 5861377
                                                                                                                                                                                                                                                                                                       Query Match 0.8%;
Best Local Similarity 62.2%;
Matches 28; Conservative
                                                                                                                                                 GENERAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEFAX: (303) 793-34
INFORMATION FOR SEQ ID NO:
                                     CORRESPONDENCE ADDRESS: ADDRESSE: No. 5861377artis Corporation
                                                                                              APPLICANT: Hans Fritz
APPLICANT: Christian Sommerhoff
APPLICANT: Jutta Heim
                                                                     TITLE OF INVENTION: NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                               FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                              FEATURE:
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NAME: Barry J. Swanson
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APPLICATION NUMBER:
FILING DATE: 07-JUNE
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APPLICATION NUMBER: 08/479,724
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                  CITY:
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MEDIUM TYPE: Diskett
                            STREET:
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CORRESPONDENCE ADDRESS:
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TOPOLOGY: 1ii
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                                                                                                                                                                                                                                                                    140 CTTCTCCAGGAAGAGCACTCAGGAGACCAGGAAAATGGCTACAGG 184
                                                                                                                                              INFORMATION:
                                                                                                                                                                                                                                           East Hanover
New Jersey
                                                                                                                                                                         Application US/08369829A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                nucleic acid
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                            59 Route 10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (303) 793-3333
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Swanson & Bratschun, L.L.C.
00 E. Prentice Avenue, Suite
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             07-JUNE-1995
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                                                                                   NO.
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                                                                                5861377el Inhibitor
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                                                                                                                                                                                                                                                                                                                     Score 21; Pred. No. 2
                                                                                                                                                                                                                                                                                                    Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Suite 200
                                                                                                                                                                                                                                                                                                                                 DB 5;
                                                                                                                                                                                                                                                                                                                  .9e+03;
                                                                                                                                                                                                                                                                                                   15; Indels
                                                                                                                                                                                                                                                                                                                               Length 59;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MB
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
Matches 25; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX: (908) 277-43
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US/08/369,829A FILING DATE: 6-JAN-95
PRIOR APPLICATION DATA:
APPLICATION NUMBER: EPO 94810006.0 FILING DATE: 7-JAN-94
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   REFERENCE/DOCKET NUMBER: 4-TELECOMMUNICATION INFORMATION: TELEPHONE: (908) 277-5110
                            PRIOR APPLICATION DATA: APPLICATION NUMBER:
                                                                                                                                                SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MOLECULE TYPE: DNA (genomic)
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                                                                                PRIOR APPLICATION DATA:
APPLICATION NUMBER: 1
                                                                                                                                                                                                                                                                                                                                                          CORRESPONDENCE
                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Fritz, Hans,
APPLICANT: Sommerhoff, Christian
TITLE OF INVENTION: Tryptase Inh
                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES:
ATTORNEY/AGENT INFORMATION
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LOCATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       REGISTRATION NUMBER:
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                                                                                                              APPLICATION NUMBER: US/08 FILING DATE: 25-JAN-1996
                                                                                                                                                                               COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE:
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                                                                FILING DATE:
                                                                                                                                                                                                                                                          COUNTRY: US
ZIP: 07901-1027
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                                                                                                                                                                                                                                                                                                            Summit
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SYSTEM: PC-DOS/MS-DOS
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DN NUMBER: 33200
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                                                                                                                                                                                                                                                                                                                                 E ADDRESS:
No. 5972698artis Corporation, Patent and Trademark Department
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Floppy disk
                                                                  25-JUL-1994
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Pred. No. 2.9e
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                                                                                                                                                                    Version #1.25 (EPO)
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                                                                                                                                                        RESULT 9
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                      Sequence 244, Applica Patent No. 6280932 GENERAL INFORMATION:
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Best Local S
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PRIOR APPLICATION NUMBER: US OPPRIOR FILTENC NUMBER: US OPPRIOR FILTENC
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CURRENT FILING DATE: 2000-08-16
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NAME/KEY: allele
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TELECOMMUNICATION INFORMATION:
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                                                             244, Application US/08952793
5. 6280932
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PARMA, et al.
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Pred. No. 3.3e+03;
1; Mismatches 15;
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Pred. No. 2.9e+03;
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8400 E.

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RESULT 10
PCT-US96-09455A-244
Sequence 244, Application PC/TUS9609455A; Sequence 11 INFORMATION:
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                                                                                                                                                                                                                                                                  Query Match
Best Local (
                                                                                                                                                                                                                                                Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEFAX: (303) 793-34 INFORMATION FOR SEQ ID NO:
                  APPLICANT: PARMA, et al.
TITLE OF INVENTION: HIGH AFFINITY NUCLEIC ACID
TITLE OF INVENTION: LICANDS TO LECTINS
NUMBER OF SEQUENCES: 390
                                                                                                                                                                                                                                                                                                                                           FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
LENGTH: 52 base pairs
                                                                                                                                                                                                                                                                                                                                                                                     MOLECULE TYPE:
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TELECOMMUNICATION INFORMATION:
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PRIOR APPLICATION DATA:
PRIOR APPLICATION NIMBER: PCT/US96/09455
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REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: 08 FILING DATE: 07-JUNE-1
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FILING DATE: 07-JUNE-1995
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8400 E. Prentice Avenue,
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Swanson & Bratschun, L.L.C
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65.78;
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                                                                                                                                                                                                                                            Mismatches
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e, Suite
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RESULT 11
US-08-374-641-40/c
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                                                                                                                                                                                                                     Patent No.
                                                                                                                                                                                                                       Sequence 40, Application US/08374641 Patent No. 5811231
                                                                                                                                                                                                                                                                                                                                                                               Best Local Similarity Matches 23; Conserv
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                                                                                                                                                                                                       GENERAL INFORMATION:
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                                                                                                                                 APPLICANT: Farr, Spencer B.
APPLICANT: Todd, Marque D
TITLE OF INVENTION: METHODS AND DIAGNOSTIC
TITLE OF INVENTION: DETERMINING TOXICITY
COMPUTER READABLE FORM:
                                                                                                       CORRESPONDENCE ADDRESS:
                                                                                                                       NUMBER OF SEQUENCES:
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TELECOMMUNICATION INFORMATION:
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NAME: Barry J. Swanson
REGISTRATION NUMBER: 33,
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                                COUNTRY:
                                                  STATE:
                                                               CITY:
                                                                            STREET:
                                                                                          ADDRESSEE:
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STRANDEDNESS: sing
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SOFTWARE: WordPer
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New York
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Colorado
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1251 Avenue of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              52 base pairs
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Pred. No. 3.
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TYPE:

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GENERAL INFORMATION:
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Best Local
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                                                            SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                  SOFTWARE: PatentIn Rel CURRENT APPLICATION DATA:
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HYPOTHETICAL:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION DATA:
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                                                                                                                       TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                     ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES:
CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION:
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Local Similarity 68.3%;
hes 28; Conservation
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                                                                                                                                                                                                    APPLICATION NUMBER: FILING DATE: Decemb CLASSIFICATION: 800
                                                                                                                                                                                                                                                                                COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
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REFERENCE/DOCKET NUMBER: X-1 CIP
             STRANDEDNESS:
                                                                                          TELEFAX:
                                                                                                        TELEPHONE:
                                                                                                                                                        NAME: Marcus-Wyner, Lynn REGISTRATION NUMBER: 34,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STRANDEDNESS:
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 TOPOLOGY:
                                                                                                                                        REFERENCE/DOCKET NUMBER:
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unknown
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                                                                                                                                                                                                                                                                                                               Floppy disk
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er 1, 1995
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Pred. No. 4.7e+03;
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; ORGANISM: oligonucleotide US-08-553-619B-13
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US-08-242-035A-2
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                                                                                                 RESULT 14
US-08-477-270-21/c
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                               Patent No. 5629158
GENERAL INFORMATION:
APPLICANT: UHLEN,
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Best Local Similarity
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                                                                   Sequence 21, Application US/08477270 Patent No. 5629158
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Best Local
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APPLICATION NIMBER

CURRENT APPLICATION DATA:
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APPLICANT: UHLEN, Mathias TITLE OF INVENTION: SOLID TITLE OF INVENTION: CONDIT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
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MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
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CORRESPONDENCE ADDRESS:
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OPERATING SYSTEM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Clark, Paul T. REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US/08/242,035A FILING DATE: May 12, 1994
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REFERENCE/DOCKET NUMBER: 00
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(617) 542-8906
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    PLASMILAR AND PLASMILAR ANTIBODY
    COMPOSITIONS AS MALIGNANT CELL MARKERS
                                                                                                                                                                                                                                                     0.7%;
   CONDITIONS
                 SOLID PHASE DIAGNOSIS OF MEDICAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         07/872,645
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1992
                                                                                                                                                                                                                                       0;
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                                                                                                                                                                                                                                  Score 20.2; DB 1;
Pred. No. 4.9e+03;
0; Mismatches 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 20.2; DB 2;
Pred. No. 4.8e+03;
                                                                                                                                                                                                                                                                        DB 1;
                                                                                                                                                                                                                                                                   Length 53;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 52;
                                                                                                                                                                                                                                       Indels
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US-08-758-626-23/c
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                                                                                                                                                                                                                                                                                                    Sequence 23, Applicat Patent No. 5736334 GENERAL INFORMATION: Dwe Sp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY: lir
MOLECULE TYPE:
DESCRIPTION:
                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy diskette
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-I
                               SOFTWARE: Wordperfect CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                  NUMBER OF SEQUENCES: 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: BENT, Stephen A.
REGISTRATION NUMBER: 29,768
REFERENCE/DOCKET NUMBER: 16
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                               TITLE OF INVENTION: I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: PatentIn Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ATTORNEY/AGENT INFORMATION:
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PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity
APPLICATION NUMBER: US/0 FILING DATE: 27-NOV-1996
                                                                                                                                                                                                 STREET:
                                                                                                                                                COUNTRY:
                                                                                                                                                                                                                  ADDRESSEE: Abbott Laboratories
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STRANDEDNESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: FILING DATE: 07-NOV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US/08/477,270 FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CENGTH:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    863 GTTTGGGCCAGGCTTTATCAAGGAGTCAAACCTCCTTAGCCTCCAGAAG 911
                                                                                                                                                                                                                                                                                                                                                                                                                                      52
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        31;
                                                                                                                                                                                   T: One Abbott Park Road Abbott Park
                                                                                                                                    60064-3500
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                                                                                                                                                               Illinois
                                                                                                                                                                                                                                                                                                                                                    Application US/08758626
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                                                                                                                                                  USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          : (703)836-9300
(703)683-4109
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Foley & Lardner
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Other nucleic acid;
Synthetic DNA oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          UMBER: US 07/781,157
07-NOV-1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       single
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63.3%;
                                                                PC-DOS/MS-DOS
                                                                                                                                                                                                                                                               NUCLEOTIDE SEQUENCES AND PROCESS FOR AMPLIFYING AND DETECTION OF HEPATITIS B VIRAL DNA
               US/08/758,626
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 20.2; DB 1;
Pred. No. 5.3e+03;
0; Mismatches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 60;
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Best Local S
Matches 26
Query Match
Best Local Similarity
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                                                                                                                                         INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                INFORMATION FOR SEQ ID NO:
                                                                                                                                                                   REFERENCE/DOCKET NUMBER: 52
TELECOMMUNICATION INFORMATION:
TELEPHONE: 708-937-4884
TELEFAX: 708-938-2623
                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM: MEDIUM TYPE: Floppy diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: TITLE OF INVENTION: NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                   ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ATTORNEY AGENT INFORMATION:
NAME: Thomas D. Brainard
REGISTRATION NUMBER: 32,459
REFERENCE/DOCKET NUMBER: 528
TELECOMMUNICATION INFORMATION:
                                                               MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
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PRIOR APPLICATION DATA:
                                                                                                          TYPE:
                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US FILING DATE: JULY 13, CLASSIFICATION:
                                                                            TOPOLOGY:
                                                                                         STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STREET: One Abbott Park Road CITY: Abbott Park
                                                                                                                                                                                                                                       REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY: USA
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                                                                                                                            LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH:
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26; Conserv
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                                                                                                          nucleic acid
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                                                                                                                                                                                                                                                   Thomas D. Brainard
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Illinois
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                                                                                                                                                                                                                                                                                                                                               Wordperfect
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                                                                          linear
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                                                           genomic DNA
                                                                                            double
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0.7%;
72.2%;
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AMPLIFYING
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                                                                                                                                                                                                                                    32,459
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                                                                                                                                                                                                                     5284.PC.01
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            5284.US.01
Score 20; DB 5;
Pred. No. 5.3e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCES AND PROCESS FOR AND DETECTION OF HEPATITIS B VIRAL DNA
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5.3e+03;
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10;

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US-08-173-968-7
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                                 Matches
                                           Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                 NFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEPHONE: (212) 527-7700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION: 514
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE CHARACTERISTICS:
LENGTH: 55 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                       PUBLICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                  MOLECULE TYPE: DI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                      FEATURE:
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                                                                                                                                                                                                                                                                                                   NTI-SENSE:
                                                                                                                                         JOURNAL: VOLUME:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: Patentin Release #1.0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             REGISTRATION NUMBER: 32,613
REFERENCE/DOCKET NUMBER: 5969/05982US1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CITY: New York
STATE: NY
                                                                                                                                                                                                                     OTHER INFORMATION: note=This sequence can be found OTHER INFORMATION: in the application, as filed, on page 25, line
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US 0 FILING DATE: 26-AUG-1991
                                                                                                           DATE:
                                                                                                                        PAGES: 14891-14894
                                                                                                                                                                        AUTHORS:
                                                                                                                                                                                                         OTHER INFORMATION:
                                                                                                                                                                                                                                                     NAME/KEY: misc_feature LOCATION: 1..55
                                                                                                                                                                                                                                                                                                                                                                STRANDEDNESS:
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618 AGTGGCAAAGCAGAATGGATTCAAGAGGGAGAAGACTCCAGACT 661
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                             l Similarity
29; Conserv
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                                                                                                           1987
                                                                                                                                                                                                                                                                                                                                                                             nucleic acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                              236687
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                             (212) 753-6237
                                                                                                                                                                        001,
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                               Conservative
                                                                                                                                                        Ooi, C. E. et al.
J. Biol. Chem.
                                                                                                                                                                                                                                                                                                     No
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                                                                                                                                                                                                                                                                                                                                                   linear
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                                                                                                                                                                                                                                                                                                                                                                single
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                                              0.78;
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                            Score 20; DB:
Pred. No. 5.8e-
0; Mismatches
                               0;
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                                                             DB 1;
                             .8e+03;
ies 15;
                                                           Length 55;
                             Indels
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RESULT 18
US-07-801-814D-7
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US-07-801-814D-7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                             TELEFAX: (212) 753-623
[NFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: Wordperfect CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                         REFERENCE/DOCKET NUMBER: 59
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CLASSIFICATION: 435 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
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TITLE OF INVENTION:
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                                                                                                                                                                           PUBLICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE:
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                         MOLECULE TYPE: HYPOTHETICAL:
                                                                                                                                                                                                                                                                                 FEATURE
                                                                                                                                                                                                                                                                                           ANTI-SENSE:
                                                                                 ISSUE: PAGES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDIUM TYPE: Diskette, 5.25 inch, MEDIUM TYPE: 360 Kb storage
                                                                                                                                                                                         OTHER INFORMATION:
OTHER INFORMATION:
                                                                                                                                                                                                                    OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                        STRANDEDNESS:
TOPOLOGY: 11:
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                                        DOCUMENT NUMBER: FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         REGISTRATION NUMBER:
          PUBLICATION DATE:
RELEVANT RESIDUES
                                                                     DATE:
                                                                                                                VOLUME:
                                                                                                                                JOURNAL:
                                                                                                                                             TITLE
                                                                                                                                                           AUTHORS:
                                                                                                                                                                                                                                  IDENTIFICATION METHOD:
                                                                                                                                                                                                                                                  LOCATION:
                                                                                                                                                                                                                                                                 NAME/KEY:
                                                                                                                                                                                                                                                                                                                                                                                   LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         , Application US/07801814D 6132775
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                                                                                                                262
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                                                                                                                                J. Biol. Chem.
                                                                                                                                                          Ooi, C.E. et al
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                                                                                                                                                                                                                                                                                                                                       linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                            (212) 527-7700
                                                                                                                                                                                                                                                                                                            No
                                                                                                                                                                                                                                                                                                                         Other nucleic acid
                                                                                                                                                                                                                                                                                                                                                      single
                                                                                                                                                                                                                                                                                                                                                                                                                             753-6237
            IN SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Peter
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BACTERICIDAL/PERMEABILITY-INCREASING
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                                                                                                                                                                                                     found
                                                                                                                                                                                                                   This sequence can
                                                                                                                                                                                      page 23;
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                                                                                                                                                                                         the application, line 7.
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Query Match

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Score 20;

DB 3;

Length 55;

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RESULT 20
US-09-258-367-21/c
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US-08-726-807B-21/c
; Sequence 21, Application US/08726807B
; Patent No. 6090618
                                                                                                                                                                                                                                                                                                                                                                      US-08-726-807B-21
                                                                                                                        GENERAL INFORMATION:
                                                                                                                                     Sequence 21, Application US/09258367
Patent No. 6114311
                                                                                                                                                                                                                                                                                                        Matches
                                                                                                                                                                                                                                                                                                                     Query Match
Best Local
                                                                  APPLICANT: PARMACEK, MICHAEL S.
APPLICANT: SOLWAY, JULIAN
TITLE OF INVENTION: PROMOTER FOR SMOOTH MUSCLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Best Local Similarity 65.9 Matches 29; Conservative
                      CURRENT APPLICATION NUMBER: US/09/258,367 CURRENT FILING DATE: 1999-02-26
                                                         FILE REFERENCE: ARCD:310
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEFAX: (512) 474-75
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS:
LENGTH: 47 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Parmacek, Michael S. APPLICANT: Solway, Julian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES: 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                      STRANDEDNESS: si
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US 6 FILING DATE: 05-OCT-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US/01
FILING DATE: 07-OCT-1996
                                                                                                                                                                                                                                                                                                                     Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CLASSIFICATION:
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                                                                                                                                                                                                                                  618 AGTGGCAAAGCAGAATGGATTCAAGAGGGGAGAAGACTCCAGACT 661
                                                                                                                                                                                                                                                                                                    21;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Houston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               McMillian, Nabeela R
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Texas
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (512) 418-3000
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91.3%;
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Pred. No. 6.1e
0; Mismatches
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0; Mismatches
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6.1e+03;
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ches 15;
                                                                   CELL EXPRESSION
                                                                                                                                                                                                                                                                                                                                 Léngth 47;
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US-09-431-414-21/c
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; OTHER INFORMATION: Primer
US-09-258-367-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-09-546-550-21
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                                SOFTWARE: Pa
SEQ'ID NO 21
LENGTH: 47
                                                                                                                                                                                                                                 GENERAL INFORMATION:
                                                                                                                                                                                                                                                  Sequence 21, Application US/09431414 Patent No. 6291211
                                                                                                                                                                                                                                                Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: PatentIn SEQ ID NO 21
                                                                        APPLICANT: SOLWAY, JULIAN
TITLE OF INVENTION: PROMOTER FOR SMOOTH MUSCLE CELL EXPRESSION
FILE REFERENCE: ARCD: 335
CURRENT APPLICATION NUMBER: US/09/431,414
CURRENT FILING DATE: 1999-11-01
EARLIER APPLICATION UNMEER: 08/726,807
EARLIER FILING DATE: 1996-10-07
NUMBER OF SEQ ID NOS: 55
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: SOLWAY, JULIAN
TITLE OF INVENTION: PROMOTER FOR SMOOTH MUSCLE CELL EXPRESSION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FEATURE:
OTHER INFORMATION:
OTHER INFORMATION:
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TYPE: DNA
ORGANISM: Artificial Sequence
ORGANISM: Artificial Sequence
                    TYPE: DNA
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5. 6284743
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                                                                  PatentIn Ver. 2.0
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91.3%;
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91.3%;
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Pred. No. 6.1e+03;
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Pred. No. 6
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OTHER INFORMATION: Description of Artificial Sequence: OTHER INFORMATION: Primer US-09-431-349C-21
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US-09-431-349C-21/c
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                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
APPLICANT: PARMACEK, MICHAEL S.
APPLICANT: SOLWAY, JULIAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
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                                                                                                                      SEQ ID NO 21
                                                                                                                                                                                                                                                                                                                                                                Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: PARMACEK, MICHAEL S.
APPLICANT: SOLWAY, JULIAN
TITLE OF INVENTION: PROMOTER FOR SMOOTH MUSCLE CELL EXPRESSION
FILE REFERENCE: ARSB:526
                                                                                                                                                                                                                                        FILE REFERENCE: ARSB:526
CURRENT APPLICATION NUMBER: US/09/431,349C
CURRENT FILING DATE: 1999-11-01
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT FILING DATE: 1999-01-05

EARLIER APPLICATION NUMBER: 08/726,807

REARLIER FILING DATE: 1996-10-07

NUMBER OF SEQ ID NOS: 55
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/09/225,670 CURRENT FILING DATE: 1999-01-05
                                                                                                                                                    PRIOR APPLICATION NUMBER: 08/726,807
PRIOR FILING DATE: 1996-10-07
NUMBER OF SEQ ID NOS: 55
                                                                                                                                                                                                        PRIOR APPLICATION NUMBER: 09/225,670 PRIOR FILING DATE: 1999-01-05
                                                                                                                                                                                                                                                                                         APPLICANT: SOLWAY, JULIAN
TITLE OF INVENTION: PROMOTER FOR SMOOTH MUSCLE CELL EXPRESSION
                                                                                                                                      SOFTWARE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: PatentIn Ver.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OTHER INFORMATION: Description OTHER INFORMATION: Primer
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: DNA
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                                                              ORGANISM: Artificial Sequence
                                                                                    TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FEATURE:
                                                  FEATURE:
                                                                                                  LENGTH:
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21; Conserv
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91.3%;
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91.3%;
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Pred. No. 6
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Pred. No. 6.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           of Artificial Sequence:
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                                      US-09-037-990B-60/c
                                                      RESULT 26
                                                                                                                                               Query Match
Best Local S
Matches 24
Sequence 60, Applicati Patent No. 6248519
GENERAL INFORMATION:
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Best Local S
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INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                     REFERENCE/DOCKET NUMBER: 08
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE: 18-JAN-1995 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                MOLECULE TYPE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER:
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                                                                                                                                                                                                                                                                                                                                                               REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE: 18-JU APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER:
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                                                                                                                                                                                                                              TOPOLOGY:
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             0, Application 6248519
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                                                                                                                                                                                                                                                                   57 base pairs
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                                                                                                                                                                                                                                                                                                            619/678-5099
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Barbas III, Carlos F. Gottesfeld, Joel M.
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                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SYSTEM: Windows 95
FastSEQ for Windows Version
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 IBM Compatible
                                                                                                                                                                                                                               linear
                                                                                                                                                                                                                                                                                                                         619/678-5070
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                                                                                                                                                                                                                                                                                                                                                                       Lisa A.,
                                                                                                                                                                                                                 Genomic DNA
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27-MAY-1997
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                           US/09037990B
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Pred. No. 6.1e+03;
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                                                                                                                                                           Score 19.8; DB Pred. No. 7e+03;
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                                                                                                                                                Mismatches
                                                                                                                                                                        DB 3; Length 57;
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APPLICANT: ENGEL,

Stacia R.

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Best Local
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INFORMATION FOR SEQ ID NO: 60:
SEQUENCE CHARACTERISTICS:
                             COMPUTER READABLE FORM:
MFDIUM TYPE: Floppy disk
                                                                                                                                                                         NERAL INCOMPANDANT APPLICANT: DORAI, HAIMANTI APPLICANT: OPPERMANN, HERMANN
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HIGH PROTEIN OF INVENTION: PRODUCTION FROM NON-NATIVE DNA
                                                                        ZIP:
                                                                                         COUNTRY:
                                                                                                                                          STREET:
                                                                                                                                                          ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity
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TELEPHONE: 312/474-6300
TELEFAX: 312/474-0448
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MEDIUM TYPE: Floppy disk
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TITLE OF INVENTION: DETECTION OF FERMENTATION-RELATED
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                                                                       07148
                                                                                                                        HOPKINTON
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: nucleic acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STRANDEDNESS: single
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US/09/037,990B FILING DATE: 11-Mar-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY: United States of America ZIP: 60606-6402
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STATE: Illinois
                                                                                                                                          E: PATENT ADMINISTRATOR, CREATIVE BIOMOLECULES, 45 SOUTH STREET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADDRESSEE: Marshall, O'Toole, Gerstein, Murray & STREET: 6300 Sears Tower, 233 South Wacker Drive
                                                                                         USA
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                  IBM PC compatible
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PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQ ID NO: 60:
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Pred. No. 7.1e+03;
0; Mismatches 17
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Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
                                                                     CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/14:
FILING DATE: 25-OCT-1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: 617/248-7100 INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                          COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
              TELECOMMUNICATION INFORMATION:
                                                                                                                                                                              CURRENT APPLICATION DATA
                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
ADDRESSEE: PATENT ADMINISTRATOR, CREATIVE BIOMOLECULES, STREET: 45 SOUTH STREET
                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: DORAI, HAIMANTI
APPLICANT: OPPERMANN, HERMANN
TITLE OF INVENTION: METHODS AN
TITLE OF INVENTION: PRODUCTION
                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: Patentin Release #1.0, Version CURRENT APPLICATION DATA:
                            REGISTRATION NUMBER: 34
REFERENCE/DOCKET NUMBER:
                                                             NAME:
                                                                                                                                                     FILING DATE:
                                                                                                                                                               APPLICATION NUMBER:
                                                                                                                                                                                                SOFTWARE:
                                                                                                                                                                                                              OPERATING SYSTEM:
                                                                                                                                                                                                                              COMPUTER:
                                                                                                                                                                                                                                                                             ZIP:
                                                                                                                                                                                                                                                                                          COUNTRY:
                                                                                                                                                                                                                                                                                                           STATE:
                                                                                                                                                                                                                                                                                                                      CITY: HOPKINTON
FELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME/KEY: misc_feature LOCATION: 1..56
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: KELLEY, ROBIN D
REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            567 GGAAAACACTATCCAGAAGATCAGAAAACAACAGC
                                                                                                                                                                                                                                                                           07148
                                                       KELLEY, ROBIN D
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                                                                                                                                                                                                                                                                                          USA
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                                                                                                                                                                                              PatentIn Release #1.0,
                                                                                                                                                                                                           IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                          Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CDNA
                                                                                                                                                  05-JUN-1995
                                                                                                                                                                                                                                                                                                                                                                                             METHODS AND
PRODUCTION
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                                                                                                      08/143,498
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                                                                                                                                                               US/08/463,675
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                                          34,637
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                             CRP093
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                                                                                                                                                                                                                                                                                                                                                                                             AND COMPOSITIONS FOR HIGH PROTEIN ION FROM NON-NATIVE DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 19.6;
Pred. No. 8
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                                                                                                                                                                                              Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        8e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            600
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US-08-464-589-9/c
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Matches 25; Conservative
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Best Local Similarity
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                                                                                                                                                                                                                              TELEFAX: 617/248-7100 INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/464,589
FILING DATE: 05-JUN-1995
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                REGISTRATION NUMBER: 34,637
REFERENCE/DOCKET NUMBER: CR
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617/248-7100
TELEPAX: 617/248-7100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US 08/143,498
FILING DATE: 25-OCT-1993
ATTORNEY/AGENT INFORMATION:
NAME: KELLEY, ROBIN D
DESCREPANTON NUMBER: 34 637
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
ADDRESSEE: PATENT ADMINISTRATOR, CREATIVE BIOMOLECULES, INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: DORAI, HAIMANTI
APPLICANT: OPPERMANN, HERMANN
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                                                                                                                             FEATURE:
                                                                                                                                            MOLECULE TYPE: cDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR HIGH PROTEIN TITLE OF INVENTION: PRODUCTION FROM NON-NATIVE DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME/KEY: misc_feature
LOCATION: 1..56
OTHER_INFORMATION: /note= "PACISIG"
                                                                      LOCATION: 1..56
OTHER INFORMATION: /note= "PAC1SIG"
                                                                                                                                                              TOPOLOGY:
                                                                                                                                                                       TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY:
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                                                                                                       NAME/KEY: misc_feature
                                                                                                                                                                                                              ENGTH:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              45 SOUTH STREET
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                                                                                                                                                                                                                56 base pairs
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0.7%;
73.5%;
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                                                                                                                                                                                                                                                                                                                       CRP093
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Pred. No. 8e+03;
Score 19.6; DB 1; Pred. No. 8e+03;
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                Length 56;
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RESULT 30
US-08-406-030A-8
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                                                 US-08-406-030A-8
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Best Local Similarity
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                                                                                                                                               INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION DATA:
APPLICATION NUMBER: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                              PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/US92/09627
FILING DATE: 05-NOV-1992
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Treco, Douglas A.
APPLICANT: Heartlein, Michael W.
APPLICANT: Hauge, Brian M.
APPLICANT: Selden, Richard F
APPLICANT: Selden, Richard F
TITLE OF INVENTION: Protein Production and Delivery
                                                                                                                                                                                                            REFERENCE/DOCKET NUMBER: TK
                                                                                                                                                                                                                                                                                                                                                                           FILING DATE: 05-NOV-
                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION DATA:
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                                                                MOLECULE TYPE:
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CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: FILING DATE: 03-DE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CITY: Lexington
                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: PCT/US93/11704 FILING DATE: 02-DEC-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: FILING DATE: 10-JUI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE:
                                                                                 TOPOLOGY:
                                                                                               STRANDEDNESS:
                                                                                                                                LENGTH:
                                                                                                                                                                                               TELEPHONE:
                                                                                                                                                                                                                                                  REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: PatentIn Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADDRESSEE: Hamilton, Brook
STREET: Two Militia Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 8, Application US/08406030A
5, 6270989
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                                                                                                             nucleic acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Massachusetts
                                                                                                                                                                              : (617) 861-6240
(617) 861-9540
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                 linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              IUMBER: US 07/911,533
10-JUL-1992
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                                                                DNA (genomic)
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                                                                                               single
0.78;
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                                                                                                                                                                                                                                               32,227
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                                                                                                                                                                                                                               TKT95-01
Score 19.6; DB 4;
Pred. No. 8.4e+03;
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             Length 60;
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Gaps

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US-08-406-030A-9
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US-08-406-030A-9/c
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Query Match
Best Local Similarity
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                                                             STRANDEDNESS:
TOPOLOGY: lin
MOLECULE TYPE:
                                                                                                                                                           NFORMATION FOR SEQ ID NO:
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                                                                                                                        SEQUENCE CHARACTERISTICS:
LENGTH: 60 base pairs
                                                                                                                                                                                        TELECOMMUNICATION INFORMATION: TELEPHONE: (617) 861-6240
                                                                                                                                                                                                                                                                                                                                 FILING DATE: 05-NOV-PRIOR APPLICATION DATA:
APPLICATION NUMBER: 1
FILING DATE: 02-DEC-
                                                                                                                                                                                                                                                                   ATTORNEY/AGENT INFORMATION:
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PRIOR APPLICATION NUMBER: US 07/985,586
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APPLICATION NUMBER: US 08/243,391
FILING DATE: 13-MAY-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                      REFERENCE/DOCKET NUMBER: TKT95-01
                                                                                                                                                                                                                                 NAME: Granahan, Patricia
REGISTRATION NUMBER: 32,227
                                                                                                                                                                                                                                                                                APPLICATION NUMBER: PCT/
FILING DATE: 05-NOV-1992
                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US/0 FILING DATE: 17-MAR-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Massachusetts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Two Militia Drive
                                                                                                                                                                        (617) 861-9540
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Hauge, Brian M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Heartlein, Michael W
                                                                                 linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Treco, Douglas A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hamilton, Brook, Smith & Reynolds, P.c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ication US/08406030A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER: US 07/911,533
10-JUL-1992
                                                                                                                                                                                                                                                                                                                                                                                                                            UMBER: US 07/787,840
05-NOV-1991
                                                                                                                                                                                                                                                                                                                                                                              05-NOV-1991
                                                             DNA (genomic)
                                                                                                                                                                                                                                                                                                                                 JMBER: PCT/US93/11704
02-DEC-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     03-DEC-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Floppy disk
                                                                                            single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Protein Production and Delivery
                                                                                                                                                                                                                                                                                                PCT/US92/09627
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Score 19.6;
Pred. No. 8
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DB 4;
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            Length 60;
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                                                                                                                                 US-08-417-210A-99
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                                                                                                                                                   RESULT 33
                                                                                                                                                                                                                                                                                                                     TOPOLOGY: US-08-169-715-14
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                                                                                 Sequence 99, Application US/08417210A Patent No. 5863542 GENERAL INFORMATION:
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US-08-169-715-14
                                                                                                                                                                                                                                                                                                                                                                    TELEFAX: 415-327-2951
TELEX: 706141
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 33 base pairs
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Best Local (
                                                                                                                                                                                                                                                             Matches
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                                     APPLICANT:
APPLICANT:
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                      TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEPHONE: 415-813-5600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE: 18-DEC-1991
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: PatentIn Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Chang, C
TITLE OF INVENTION:
TITLE OF INVENTION:
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                                                                                                                                                                                                                                                                        Local
                                                                                                                                                                                                                                                                                                                                                       STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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REFERENCE/DOCKET NUMBER: 22300-20150.00
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE: Morrison & Foerster STREET: 545 Middlefield Road, Suite 200 CITY: Menlo Park
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER:
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                                                                                                                                                                                                                                                             20;
                                                                                                                                                                                             1 TCTCCAYTTRGTRCTGTCYTTTTTCTTTATR 31
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                                                                                                                                                                                                                                                                        0.7%;
Similarity 64.5%;
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                                                                                                                                                                                                                                                                                                                                                                 : 33 base pairs nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Monroy, Gladys H
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Application US/08169715
                                   TARTAGLIA, JAMES COX, WILLIAM I.
                                                                    PAOLETTI, ENZO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Chang, Chu-An
                                                                                                                                                                                                                                                           Conservative
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                                                                                                                                                                                                                                                                                                                                         linear
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                                                                                                                                                                                                                                                                                                                                                      single
                   IMMUNODEFICIENCY RECOMBINANT POXVIRUS
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        148
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     HIV PROBES FOR USE IN SOLUTION PHASE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         07/813,583
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                                                                                                                                                                                                                                                                                                                                                                                                                   14:
                                                                                                                                                                                                                                                        Score 19.4; DB 4;
Pred. No. 6.5e+03;
5; Mismatches 6;
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                                                                                                                                                                                                                                                                                     Length 33;
                                                                                                                                                                                                                                                      Indels
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Gaps

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STREET: 530 FIFTH AVENUE
CITY: NEW YORK
STATE: NEW YORK

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Query Match
Best Local Similarity
"~+~hes 26; Conserv:
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                                                                                                                                                                  SOFTWARE: Pa
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 760, Application US/09641638
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Patent No. 6432648
          -09-641-638-760
                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: US 60/133,200 PRIOR FILING DATE: 1999-05-07 PRIOR APPLICATION NUMBER: US 09/275,267 PRIOR FILING DATE: 1999-03-23 PRIOR APPLICATION NUMBER: US 60/119,917 PRIOR FILING DATE: 1999-02-12
                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/09/641,638
CURRENT FILING DATE: 2000-08-16
PRIOR APPLICATION NUMBER: US 09/502,330
PRIOR FILING DATE: 2000-02-11
                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: BIALLELIC MARKERS DERIVED FROM GENOMIC REGIONS CARRYING TITLE OF INVENTION: GENES INVOLVED IN ARACHIDONIC ACID METABOLISM FILE REFERENCE: GENSET.051CP1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT:
FEATURE:
NAME/KEY: allele
LOCATION: 24
OTHER INFORMATION: 12-400-217: polymorphic base A or
                                                                                                                                                                                                        NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   INFORMATION FOR SEQ ID NO:
                                                                                                     ORGANISM: Homo Sapiens
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MEDIUM TYPE: Floppy disk
                                                                                                                                                 ENGTH: 47
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: KOWALSKI, THOMAS J.
REGISTRATION NUMBER: 32,147
REFERENCE/DOCKET NUMBER: 454310-2690
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US/08/417,210A FILING DATE: 05-APR-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEPHONE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2223 TGTGTGTGATTATGCATGAGACTGTACTGGTAAGACT 2259
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Cohen, Annick
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    USA
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70.3%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     11;
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US-08-417-210A-106/c
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US-08-973-965-1/c
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                                                                                                                          GENERAL INFORMATION:

APPLICANT: HIMMLER, Gottfried

APPLICANT: SCHLEDERER, Thomas
                                                                                                                                                                           Sequence 1, Application US/08973965
Patent No. 6033881
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 106, Application US/08417210 \rho Patent No. 5863542
                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local (
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CURRENT APPLICATION NUMBER: US/08/973,965
CURRENT FILING DATE: 1998-04-06
EARLIER APPLICATION NUMBER: AT A 1007/95
EARLIER FILING DATE: 1995-06-13
                                                                       TITLE OF INVENTION: PROCESS FOR THE TRANSCRIPTIONLESS AMPLIFICATION OF TITLE OF INVENTION: NUCLEIC ACIDS FILE REFERENCE: 030560-055 .
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY: 1: MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELECOMMUNICATION INFORMATION: TELEPHONE: 212-840-3333
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: TARTAGLIA, JAMES APPLICANT: COX, WILLIAM I TITLE OF INVENTION: IMMUNOD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CITY: NEW YORK
STATE: NEW YORK
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME: KOWALSKI, THOMAS J. REGISTRATION NUMBER: 32,147
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: FILING DATE: 05-AP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: PatentIn Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM PC OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MEDIUM TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STREET:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1460 TCACTTTAGCTGGAAATCAAACCTCAAAACACACCAGAGGACACACT 1506
                                                                                                                                                                                                                                                                                                                            2223 TGTGTGTGATTATGCATGAGACTGTACTGGTAAGACT 2259
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              47 TGACTTTAGTTGAAAAATGTACCYAGTATTTCAGTAGAAACTCACT 1
                                                                                                                                                                                                                                                                                           40 TTTGTATCGTAATGATTGAGACTGTACCAGTAAAATT 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  . 10036
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                                                                                                                                                                                                                                                                                                                                                                                   Similarity
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                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            linear
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IBM PC compatible
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61.7%;
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                                                                                                                                                                                                                                                                                                                                                                0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                 Score 19.4; DB 2;
Pred. No. 9.3e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         454310-2690
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 19.4; DB 4;
Pred. No. 8.2e+03;
                                                                                                                                                                                                                                                                                                                                                                                                    Length 56;
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                                                                    Cuery Match
Best Local
                                                         Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Patent No. 6150087
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                                                                                                                                                                                                                TELEFAX: (508)359-3885
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   EARLIER APPLICATION NUMBER: PCT/AT96/00106
EARLIER FILING DATE: 1996-06-13
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TYPE: DNA
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                                                                                                                                                  MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/0:
FILING DATE: 14-MAR-1995
                                                                                                                                                                                                                                                          TELEPHONE: (508)359-3876
                                                                                                                                                                                                                                                                                                                          ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: Patentin Rel
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
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                                                                                                                        DESCRIPTION:
                                                                                                                                    DESCRIPTION:
                                                                                                                                                                       STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version
                                                                                                                                                               TOPOLOGY:
                                                                                                                                                                                                                                                         TELEPHONE:
                                                                                                                                                                                                                                                                                     REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                   REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER:
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                                                                                                                                                                                                      ENGTH:
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                                                       l Similarity
27; Conserv
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                                                                                                                                                                                       nucleic acid
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                         TGGAAATTCCATCTTCTAAACCAAAACTCATTGCTCAGCT 370
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TGGGAATTCCATAATGAGACCCTTAAGGTATTACTCAGCT 44
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Emeryville
                                                                                                                                                                                                                                                                                                           Harbin, Alisa A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CA
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                                                       Conservative
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                                                                                                                                                              linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Chien, David Y
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Chiron Corporation
                                                                                                                     other nucleic acid /desc = "adaptor in an expression cassette."
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                                                                 0.78;
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                                                                                                                                                                                                                                                                                             33,895
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                                                              Score 19.2; DB 3;
Pred. No. 9.2e+03;
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                                                   Mismatches
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                                                                              DB 3;
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                                                                           Length 44;
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                                                   Indels
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US-08-171-389-100
                                                           Matches
                                                                      Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 100, Patent No. 55
                                                                                                                                                                                                                                                                               TELEFAX: (415) 324-09 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
                                                                                                                                                               ORIGINAL SOURCE:
                                                                                                                                                                       MOLECULE TYPE: DI
HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                           TELECOMMUNICATION INFORMATION: TELEPHONE: (415) 324-0880
                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE: 22-JUN-1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US 0 FILING DATE: 27-JUN-1991 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS:
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APPLICATION NUMBER: US 0
FILING DATE: 17-SEP-1993
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APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                        REGISTRATION NUMBER: 33, REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                          TOPOLOGY:
                                                                                                                                                                                                                      STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       OPERATING SYSTEM:
SOFTWARE: Patentl
                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDIUM TYPE:
                                                                                                                                                                                                                                                   ENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY:
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                        1353 TACATTTGCAGGGAGTGTGAGCAAGGCTTTAG 1384
3 TAAATTAGCAGGGAGTGTTATAAAAACTTTGG 34
                                                                                                                                                                                                                               nucleic acid
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                                                                                                                                                                                                                                                                                                                                                                     Fabian, Gary R.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         5578444
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                                                          Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Andrews,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Cantor, Charles
                                                                                                                                                                                                     linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Turin,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Edwards,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                        DNA (genomic)
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                                                                                                                                                                                                                                                                                                                                                                                                JMBER: US 08/081,070
22-JUN-1993
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23-DEC-1992
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                                                                                                                                                                                                                                                                                           324-0960
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                                                                      0.7%;
75.0%;
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                                                                                                                                Human CYP2C8 gene for cytochrome P-450
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                                                                     Score 19.2;
Pred. No. 9
                                                        Mismatches
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                                                                   3e+03
                                                                                  DB 1;
                                                       8
                                                                                 Length 45;
                                                       Indels
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RESULT 39 US-08-123-936-100

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Вb
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                                                                                                                            Patent No.
                                                                                                                              Sequence 100, Application US/08475228A Patent No. 5869241
                                                                                                                                                                                                                                                                                          Matches
                                                                                                                                                                                                                                                                                                                       Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
APPLICANT: Edward
                                                                                                             GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEFAX: (415) 324-09 INFORMATION FOR SEQ ID NO:
                                                                            APPLICANT: APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                   MOLECULE TYPE: DN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
LENGTH: 45 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           REGISTRATION NUMBER: 33,875
REFERENCE/DOCKET NUMBER: 4600-0075.32/G19P2
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/723,618
FILING DATE: 27-JUN-1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/996,783
FILING DATE: 23-DEC-1992
                               APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                     HYPOTHETICAL: NO ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION:
                                                                APPLICANT:
                                                                                                                                                                                                                                                                                        Local Similarity hes 24; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                      INDIVIDUAL ISOLATE: INDIVIDUAL ISOLATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MEDIUM TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADDRESSEE:
                                                                                                                                                                                                                                                        1353 TACATTTGCAGGGAGTGTGAGCAAGGCTTTAG 1384
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       94063
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                   INVENTION
                                                                                                                                                                                                                           TAAATTAGCAGGGAGTGTTATAAAAACTTTGG 34
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CA
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                             Andrews, Beth M.
Turin, Lisa M.
Fry, Kirk E.
                                                                                              Edwards,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Edwards, Cynthia A. Cantor, Charles R.
                                                                                 Cantor,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      USA
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                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Andrews,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   E: Floppy disk
IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                      linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (415) 324-0880
                                                                                                                                                                                                                                                                                                                                                                                                                                  DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gary R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               324-0960
                                                                             Charles R.
                                                                                                                                                                                                                                                                                                     0.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DNA-Binding Molecules
Sequence-Directed DNA Binding Molecules, Compositions and Methods
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Beth M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Screening Assay for the Detection
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 640
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                                                                                                                                                                                                                                                                                                                                                                      Human CYP2C8 gene for cytochrome P-450
                                                                                                                                                                                                                                                                                          0,
                                                                                                                                                                                                                                                                                                       Score 19.2; DB 1;
Pred. No. 9.3e+03;
                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Version #1.25
                                                                                                                                                                                                                                                                                                                      Length 45;
                                                                                                                                                                                                                                                                                          Indels
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US-08-482-080A-100
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                                                                                                                                                                                  Patent No.
                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
                                                                                                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                       Matches
                                                                                                                                                                GENERAL INFORMATION:
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(NFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                REFERENCE/DOCKET NUMBER: 46
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORIGINAL SOURCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US 01
FILING DATE: 22-JUN-1993
ATTORNEY/AGENT INFORMATION:
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                           CORRESPONDENCE ADDRESS:
                                                           TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                               HYPOTHETICAL:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MOLECULE TYPE:
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                                                                                        APPLICANT:
                                                                                                                                     APPLICANT:
                                         NUMBER OF SEQUENCES:
                                                                                                      APPLICANT:
                                                                                                                      APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US 07/723,618
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STRANDEDNESS:
            ADDRESSEE
                                                                                                                                                                                                                                                                                                                                                                                                                 INDIVIDUAL ISOLATE:
                                                                                                                                                                                                                                                                                             1353 TACATITGCAGGGAGTGTGAGCAAGGCTTTAG 1384
                                                                                                                                                                                100, Application US/08482080A
5. 6010849
                                                                                                                                                                                                                                                                                                                                       24;
                                                                                                                                                                                                                                                                            3 TAAATTAGCAGGGAGTGTTATAAAAACTTTGG 34
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     T: 505 Penobscot Drive Redwood City
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            94063
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Stratford, Carol A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        45 base pairs
505 Penobscot Drive
                                                                                     Fry, Kirk E
                                                                                                                 Edwards, Cynthia A. Cantor, Charles R. Andrews, Beth M.
                                                                                                     Turin,
                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              linear
         Genelabs Technologies, Inc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Genelabs Technologies, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (415) 324-0880
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             27-JUN-1991
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06-JUN-1995
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           double
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                                                                                                      Lisa M.
                                                        Sequence-Directed DNA Binding Molecules, Compositions and Methods
                                                                                                                                                                                                                                                                                                                                                     0.7%;
                                           664
                                                                                                                                                                                                                                                                                                                                                                                                                 Human CYP2C8 gene for cytochrome P-450
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                                                                                                                                                                                                                                                                                                                                                    Score 19.2;
Pred. No. 9.
                                                                                                                                                                                                                                                                                                                                    Mismatches
                                                                                                                                                                                                                                                                                                                                                       .3e+03
                                                                                                                                                                                                                                                                                                                                                                    DB 2;
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RESULT 42
US-09-354-947-100
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                                                                                                                                                                                               Patent No.
                                                                                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                               Query Match
                                                                                                                                                                                GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEFAX: (650) 324-0960 INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                     APPLICANT: Turin, Li.
APPLICANT: Fry, Kirk
TITLE OF INVENTION:
TITLE OF INVENTION:
                                         CORRESPONDENCE ADDRESS:
                                                                                                                                   APPLICANT: Edwards, Cynthia A
APPLICANT: Cantor, Charles R.
APPLICANT: Andrews, Beth M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MOLECULE TYPE: DN.
                                                          NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                          HYPOTHETICAL: 1
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TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US 01
FILING DATE: 22-JUN-1993
ATTORNEY/AGENT INFORMATION:
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PRIOR APPLICATION UNBER: US 07/723,618
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APPLICATION NUMBER: US 07/996,783
FILING DATE: 23-DEC-1992
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PRIOR APPLICATION NUMBER: US 08/123,936
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEPHONE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US 08 FILING DATE: 17-SEP-1993
                              ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY:
                                                                                                                                                                                                                                                                                                                1353 TACATTTGCAGGGAGTGTGAGCAAGGCTTTAG 1384
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5. 6384208
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                                                                                                                                                                                                                                                                                                                                                                   Similarity
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Redwood City
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            E: Genelabs Technologies, Inc. 505 Penobscot Drive
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                                                                                                                                                                                                                                                                                                                                                   Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       linear
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, John F.
39,118
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   double
                                                                                                     Kirk E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      20-DEC-1993
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                                                                                                                    Lisa M.
                                                        Sequence-Directed DNA Binding Molecules, Compositions and M 664
                                                                                                                                                                                                                                                                                                                                                                0.7%;
                                                                                                                                                                  Cynthia A.
                                                                                                                                                                                                                                                                                                                                                                                                                            Human CYP2C8 gene for cytochrome P-450
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                                                                                                                                                                                                                                                                                                                                                              Score 19.2; DB 3;
Pred. No. 9.3e+03;
                                                                     Compositions and Methods
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                                                                                                                                                                                                                                                                                                                                                 8
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                                                                                                                                                                                                                                                                                                                                              Gaps
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RESULT 43
PCT-US93-12388-100
; Sequence 100, Application PC/TUS9312388
; GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                             Best Local Similarity
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COMPUTER READABLE FORM
                                                                                             CORRESPONDENCE ADDRESS:
ADDRESSEE: Genelabs
                                                                                                                                             TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              HYPOTHETICAL: NO ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MOLECULE TYPE: UN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: Brady, John F.
REGISTRATION NUMBER: 39,118
REFERENCE/DOCKET NUMBER: 46
TELECOMMUNICATION INFORMATION:
TELECHONE: (650) 324-0880
                                                                                                                              NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US 0
FILING DATE: 22-JUN-1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
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                              COUNTRY:
                                                               CITY: Redwood City
                                                                                STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: nucleic acid STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US 07/723,618 FILING DATE: 27-JUN-1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: FILING DATE: 17-SE
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                 94063
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                                               CA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          45 base pairs
                                USA
                                                                              505 Penobscot Drive
                                                                                                                                                                                                                                                                                                                                                               Conservative
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                                                                                            Genelabs Technologies,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            linear
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23-DEC-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             JMBER: US 08/081,070
22-JUN-1993
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07-JUN-1995
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                                                                                                                              Sequence-Directed DNA Binding Molecules, Compositions and M 641
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75.0%;
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Pred. No. 9.3e
0; Mismatches
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US-08-859-649-10
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             REFERENCE/DOCKET NUMBER: 46
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 324 0880
                            SOFTWARE: PatentIn Release #1.0, Version CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/859,649 FILING DATE:
                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: LBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOCTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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HYPOTHETICAL: NO
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PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                      NUMBER OF SEQUENCES: 3
                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT:
                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: Antibodies
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STRANDEDNESS: double
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                                                                                                                                                                                                    CITY: Menlo Park
STATE: California
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REGISTRATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: PCT/US93/12388
                                                                                                                                                                                  COUNTRY:
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5. 5874060
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                                                                                                                                                                                                                                      2730 Sand Hill Road
                                                                                                                                                                                                                                                                                                                                                                                            Armour, Kathryn L. Carr, Francis J.
                                                                                                                                                                                                                                                                                                                      Garin-Chesa, Pilar
VENTION: Recombinant Human Anti-Lewis
                                                                                                                                                                                                                                                                                                                                                                          Old, Lloyd J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                          Kitamura, Kunio
                                                                                                                                                                                    U.S.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US 08/123,936
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Pred. No. 9.3e+03
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US-08-207-861-10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 48 base ~~ Tyder
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
-08-207-861-10
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                                                                                                   INFORMATION FOR SEQ ID NO:
                                                                                                                                           REFERENCE/DOCKET NUMBER: 76
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-854-3660
                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELECOMMUNICATION INFORMATION: TELEPHONE: 415-854-3660
                                                                     SEQUENCE CHARACTERISTICS:
LENGTH: 48 base pairs
                                                                                                                                                                                                                                                                                 CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MOLECULE TYPE:
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                                                                                                                                                                                                                    CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION:
                                                                                                           TELEFAX: 415-854-50
TELEX: 66141 PENNIE
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OPERATING SYSTEM:
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mes 27; Conserva
                          TOPOLOGY:
                                                                                                                                                                                          NAME: Halluin, Albert REGISTRATION NUMBER: 2
                                                                                                                                                                                                                                                                                                SOFTWARE: PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                       STATE:
                                                                                                                                                                                                                                                                                                                                                                                                                      CITY: Menlo Park
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                                         STRANDEDNESS:
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                                         nucleic acid
EDNESS: unknown
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                                                                    48 base pairs
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                                                                                                                                415-854-3694
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Old, Lloyd J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Armour, Kathryn L.
                           unknown
                                                                                                                                                                                                                                                                                                              IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    Pennie &
                                                                                                                                                                                                                                                  08-MAR-1994
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0.7%;
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Pred. No. 9.7e+03;
0; Mismatches 13;
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US-08-431-527A-6
                          Sequence 53, Application US/08485355B
Patent No. 6177075
Patent INFORMATION:
APPLICANT: Christian, P. D., Gordon, K.
TITLE OF INVENTION: Insect Viruses and
                                                                                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                                            Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 6, Application US/08431527A Patent No. 5843650
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SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: Twinhead Slimnote-890TX
OPERATING SYSTEM: WS DOS version 6.2,
OPERATING SYSTEM: Windows version 3.11
SOFTWARE: Word for Windows version 2.0
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                     REFERENCE/DOCKET NUMBER: 12
TELECOMMUNICATION INFORMATION:
TELEPHONE: 972-3-6938541
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION DATA: APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES: 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT:
                                                                                                                                                                                                                                                             Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDIUM TYPE: 1.44 megabyte, 3.5" microdisk COMPUTER: Twinhead Slimnote-890TX OPERATING SYSTEM: MS DOS version 6.2,
                                                                                                                                                                                                                                                                                                                                            STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: Friedmam, Mark REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                             TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                        TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                          TELEX:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STREET:
                                                                                                                                                                                                                                                                                                                                                                          LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADDRESSEE:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    472 TTTTCTCCAGTCAGCAAGCTCTCAGCCAACATGTGTGGCT 511
                                                                                                                                                                                                 27;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1: 2940 Birchtree space lane
Silver Spring
Maryland
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity
                                                                                                                                                                             TACATGTGTAACAGTTCCTGCATGGGCGGCATGAACCGGAGGCCCCATC 48
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TTGGCCCCAGTAAGCAAACCACGAGCCATCACGGGTGCCT 40
                                                                                                                                                                                                                                                                                                                                                                                                                                        972-3-6938542
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          David Segev
VENTION: NO.
                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               United States of America
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                             linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mark M. Friedman c/o Robert Sheinbein
                                                                                                                                                                                                                                                                                                                                         single
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                                                                                                                                                                                                                                                          0.7%;
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                Protecting Plants
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    33,883
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                                                                                                                                                                                                                                           Score 19.2; DB 2;
Pred. No. 1.1e+04;
0; Mismatches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 19.2; DB 4;
Pred. No. 9.7e+03;
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                                                                                                                                                                                                                                                                           DB 2;
                               . H.J., Hanzlik,
Their Uses in
                                                                                                                                                                                                                                             18;
                                                                                                                                                                                                                                                                       Length 58,
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                                                                                                                                                                                                                                           Indels
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                                                                                                                                                                                                                                                                                                                        Sequence
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RESULT 48
US-08-373-124A-185/c
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SEQUENCE DESCRIPTION: SEQ ID NO: 53:

US-08-485-358-53
                                                                                                                                                                                                                                                                                                                                                                          Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEX: 910 277299
INFORMATION FOR SEQ ID NO: 53:
SEQUENCE CHARACTERISTICS:
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette,
MEDIUM TYPE: storage
                                                                                                                                                                                                                                           TITLE OF
                                                                                                                                                                                      CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                            TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                APPLICANT:
                                                                                                                                                                                                                                                                                                 APPLICANT:
                                                                                                                                                                                                                                                                                                                      APPLICANT:
                                                                                                                                                                                                                                                                                                                                     APPLICANT:
                                                                                                                                                                                                         NUMBER OF SEQUENCES:
                                                                                                                              STREET:
                                                           ZIP:
                                                                        COUNTRY:
                                                                                               STATE:
                                                                                                                                                                    ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELECOMMUNICATION INFORMATION: TELEPHONE: (415) 781-1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      759 GTGGATATAGGGTCCAGCCCTGAACGGAGGGCAGATCTAGAGGAAACA 806
                                                                                                                                                                                                                                                                                                                                                          INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 GGGGATCCACAGTTCTGCCTCCCCCGGACGGTAAATATAGGGGAACCA 48
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          l Similarity
30; Conserv
                                                         90071
                                                                                        Los Angeles
California
                                                                                                                                                                                                                                         NT: McSwiggen, James
NT: Jarvis, Thale
F INVENTION: METHODS A
                                                                                                                                                                                                                                                                                                                                                                      85, Application US/08373124A
5646042
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US 08/440,522
FILING DATE: 12-MAY-1995
APPLICATION NUMBER: US 08/089,372
FILING DATE: 08-JUL-1993
APPLICATION NUMBER: AU PL4081/92
FILING DATE: 14-AUG-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US/08/485,355B FILING DATE: 07-Jun-1995 CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STRANDEDNESS: unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: Trecartin, Richard F. REGISTRATION NUMBER: 31,801
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADDRESSEE: Flehr Hohbach Test Albritton & Herbert LLP STREET: Four Embarcadero Center, Suite 3400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX: (415)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           REFERENCE/DOCKET NUMBER: A-58631-2/RFT/DSS
                                                                                                                                                   633 West Fifth Street
                                                                          U.S.A.
                                                                                                                                Suite 4700
                                                                                                                                                                                                                                                                                                                  Draper, Kenneth
                                                                                                                                                                                                                                                                                                                                   Stinchcomb, Dan T.
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                                                                                                                                                                  Lyon & Lyon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   United States
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     unknown
                                                                                                                                                                                                                          METHODS AND COMPOSITIONS FOR TREATMENT OF RESTENOSIS AND CANCER USING RIBOZYMES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       398-3249
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Pred. No. 1.1e+04;
0; Mismatches 18;
                     1.44 Mb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 59;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0
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OPERATING SYSTEM:

IBM Compatible

IBM P.C.

DOS 5.0

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Вþ
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US-08-435-628-185/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 185, Application US/08435628 Patent No. 5817796
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT:
APPLICANT:
APPLICANT:
                                                                     OPERATING SYSTEM: IBM P.C. DOS 5.0 SOFTWARE: Word Perfect 5.1 CURRENT APPLICATION DATA:
                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette,
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELECOMMUNICATION INFORMATION: TELEPHONE: (213) 489-1600 TELEFAX: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE: May 18,
                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR TITLE OF INVENTION: TREATMENT OF RESTENOSIS AND TITLE OF INVENTION: CANCER USING RIBOZYMES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS:
LENGTH: 38 base pairs
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                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                         ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
CITY: Los Angeles
STATE: California
                            APPLICATION NUMBER: FILING DATE: 05-MA' CLASSIFICATION: 51
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nes 25; Conserv
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STRANDEDNESS: single
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TELEX: 67-3510
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE: May 18, 1994
APPLICATION NUMBER: 08/192,943
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE:
                                                                                                                               COMPUTER:
                                                                                                                                                                                                          COUNTRY:
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APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           38
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                                                                                                                                                                                            90071
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                                                                                                                                                                                                          U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                       McSwiggen, James
Jarvis, Thale
                                                                                                                                                                                                                                                                                                                                                                                                                         Draper, Kenneth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Word Perfect 5.1
                                                                                                                          IBM Compatible
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December 7, 1992
                                             05-MAY-1995
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                                                             US/08/435,628
08/373,124
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Pred. No. 9
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                                                                                                                                                            1.44 Mb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      .6e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0,
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; TYPE: DNA; ORGANISM: HUMAN US-09-609-816-13
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В
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                                                                 Query Match
Best Local Similarity
Matches 28; Conserv
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Sequence 13, Apri-
Sequence 13, Apri-
0. 6436684
                                                                                                                                                                                                    SEQ ID NO 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: ISOLATED HUMAN DRUG-METABOLIZING TITLE OF INVENTION: PROTEINS, NUCLEIC ACID MOLECULES ENCODING HUMAN TITLE OF INVENTION: DRUG-METABOLIZING PROTEINS, AND USES THEREOF FILE REFERENCE: CL000669PCT
                                                                                                                                                                                                                  NUMBER OF SEQ ID NOS: 23 SOFTWARE: FastSEQ for Windows Version
                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: 60/192,408 PRIOR FILING DATE: 2000-03-27 PRIOR APPLICATION NUMBER: 60/212,725
                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/09/609,816
CURRENT FILING DATE: 2000-07-03
                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT:
                                                                                                                                                                                                                                                                    PRIOR FILING DATE: 2000-06-20
PRIOR APPLICATION NUMBER: 09/609,816
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Woodage,
                                                                                                                                                                                                                                                      PRIOR FILING DATE: 2000-07-03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                    LENGTH: 46
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
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TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
TELEX: 67-3510
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE: August 26, ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: Warburg, Richard REGISTRATION NUMBER: 3
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TOPOLOGY: linear
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| 44434<br>44434<br>4444<br>4443<br>4443   | 430<br>431<br>432               | 426 2 428 2 428   | 422                                  | 420                           | C 416<br>C 417<br>C 418       | c 414<br>415        | c 412<br>c 413                | 409<br>410           | 407<br>C 408                | 405                          | c 402               |                     | c 398                | c 396               | 394<br>395         |                     | c 390                | 388                | 0 387               |                   | C 383              | C 381         | 379                | c 370                | 374                  | 37:<br>C 37: |
|--|---------------------------------|---|--------------------------------------|-------------------------------|-------------------------------|---------------------|-------------------------------|----------------------|-----------------------------|------------------------------|---------------------|---------------------|----------------------|---------------------|--------------------|---------------------|----------------------|--------------------|---------------------|-------------------|--------------------|---------------|--------------------|----------------------|----------------------|--------------|
| 17.6<br>17.6<br>17.6<br>17.6<br>17.6<br>17.6<br>17.6<br>17.6   | 17.6<br>17.6                    | 17.6<br>17.6<br>17.6  | 17.6<br>17.6<br>17.6                 | 17.6<br>17.6                  | 17.8<br>17.8<br>17.8          | 17.8<br>17.8        | 17<br>17                      | 17<br>17             | 17                          | 17<br>17                     | 17.<br>17.          | <u> </u>            | 17.<br>17.           | 17.                 | 17.<br>17.         | 17.<br>17.          | 17.<br>17.           | 17.                | 17.                 | 17.               | 17.                | 17.           | 17.                | 17.8                 | 17.<br>17.           | 17.8         |
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| 46 10<br>46 10<br>46 17<br>47 17<br>47 17<br>48 9<br>48 9<br>48 9<br>48 9<br>17  |                                 | 40 9<br>40 9<br>42 10<br>42 17  |                                      |                               |                               |                     | سر سر ب                       |                      |                             |                              | ب ب                 |                     |                      | 555<br>91           |                    |                     |                      |                    |                     |                   |                    |               |                    |                      |                      |              |
| HSW<br>0 AV<br>4 N4<br>7 AZ<br>7 AZ<br>7 AZ<br>7 AZ<br>7 AZ<br>8U2<br>8U2<br>8U2<br>8U2<br>8U2<br>8U2<br>8U2<br>8U2<br>8U2   | 7 AZ<br>7 AZ<br>AU2             | AAS<br>AIS<br>O AV  | AUC<br>AUC<br>AJZ<br>7 AZ            | 7 TI                          | 7 7 7<br>CC 8                 | 7 7 E               | NO 7                          | 2 0<br>B             | .7 A.                       | . 3. 7<br>: <sub>В</sub> . В |                     |                     | 74<br>AF             | 17 T                |                    |                     | 17 A                 | 10 A<br>17 A       | 17 A<br>17 T        | 9 AU              | 17 A               | AI P          | 14 AA              | 9 AU                 | 9 AC                 | 9<br>AC      |
| HSM010239<br>AV850103<br>N44913<br>AZ496495<br>D12221<br>AZ312766<br>AUZ64490<br>AUZ64490<br>AUZ64985<br>AUZ64985<br>AUZ64985<br>AUZ64985<br>AUZ64985<br>AUZ64985<br>AUZ64985<br>AUZ64985<br>AUZ64985<br>AUZ64985  | 841825<br>65123                 | 73004<br>65182<br>7952183<br>647857   | )12250<br>)12255<br>!39975           | 1108D08Q<br>258235            | 1630370<br>VS04I4H<br>VS04KW9 | 2322796             | 2484130<br>2634679<br>2633132 | V531442<br>F535695   | 43162<br>2320565<br>4862615 | H758527<br>J052665           | Z618308 .<br>03160  | 911978<br>255523    | 34759<br>2328238     | A170H10Q            | Z326602<br>H809196 | W444382<br>51371    | Z665895<br>890674    | W245427<br>Z421485 | L754632<br>A243A09P | 061719            | E636997<br>Z589567 | 040698        | 138783             | 107322               | 1105769              | 1105414      |
| · ·  |                                 |   |                                      |                               |                               |                     |                               |                      |                             |                              |                     |                     |                      |                     |                    |                     |                      | •                  |                     |                   |                    |               |                    |                      |                      |              |
| AUJS6834 AUZ65834 A1045389 Homo sapi AV850103 AV850103 N44913 YY21h06 r1 AZ496495 IM00332N24 D12221 HUM0005362 AZ312766 IM0028K12 AUZ64490 AUZ64490 AUZ64492 AUZ64490 AUZ64492 AUZ64492 AUZ64985 AUZ64985 AUZ69985 AUZ69329 AZ401474 IM0188002 AZ940011 ZM0199C14  | AZ779<br>AZ841<br>AU2651        | AA973<br>AI5651<br>AV952<br>AZ647   | AU0123<br>AU0123<br>AJ2399           | AL460<br>AU2582               | BH63(<br>AL29)                | AZ32:               | AZ48                          | AV53                 | H451<br>AZ32                | вн75<br>вл05                 | AZ61<br>B031        | AA911<br>AN915      | F347                 | AL47                | AZ32               | AW44                | AZ66                 | AW24               | AL75<br>AL48        | AL76<br>AU061     | BF63<br>AZ58       | B001<br>AI040 | AA551<br>H387      | AU107<br>AU107       | AU105<br>AU105       | AU105        |
| 34 AU2<br>39 HOM<br>103 AV<br>3 YY21<br>1495 IM<br>11 HUMO<br>1766 IM<br>90 AU2<br>90 AU2<br>92 AU2<br>85 AU2<br>85 AU2<br>474 IM<br>474 IM  | 9401 2N<br>825 2N<br>23 AU2     | )04 op:<br>182 tq:<br>183 AV  | 250 AU<br>255 AU<br>975 AJ<br>376 2N | 0901 T                        | 0370 10<br>1770 Te            | 2796 11<br>6534 11  | 4130 1<br>4679 u              | 1442 A               | 62 yo0<br>0565 11           | 8527 S<br>2665 B             | 8308 1<br>60 csr    | 978 og<br>523 All   | 59 HSP               | 9196 1<br>4190 T    | 6602 1             | 4382 A              | 5895 1               | 5427 2             | 4632 A              | 3402 A<br>719 AU  | 16997 N<br>19567 1 | .69 csF       | .834 nj<br>183 yp4 | 1322 AU<br>1633 AU   | 309 AL               | 6414 AL      |
| 265834<br>50103.<br>1005.rd<br>100332N2<br>10028K1<br>10028K1<br>164490<br>164492<br>164985<br>169329<br>1016800   | 40015 H1<br>4014 0 J1<br>265123 | 31d03.s<br>56h10.x<br>7952133   | 012250<br>012255<br>039975           | . bruce<br>258235             | 0070831<br>etraode            | M0043G              | M031012                       | V53144               | 1d10.s                      | ALK_02<br>J05266             | M0449P<br>L-169H    | 21e09.              | D30061               | . bruc              | M0049N             | B509 P              | M0547F               | 822975<br>M01996   | rabido<br>bruc      | rabidc<br>1061719 | IF074AC<br>M0398F  | :18d06.       | 58d08.             | 1107322              | 1105622<br>1105769   | 1105414      |
| 4 0 04   | ίν α                            | 4   | <b>A</b>                             | 1                             | 3 5 6                         | [4<br>[4            | ź×ω                           |                      | 301                         | 50                           | 6 2 2               | 8 4                 | >                    | e 6                 | 02                 | t.                  | 01                   | λω!                | ps<br>ei            | ps                | 08<br>2<br>1       | 10<br>S       | 1 S                |                      |                      |              |
| ·  |                                 |   |                                      |                               |                               |                     |                               |                      |                             |                              |                     |                     |                      |                     |                    |                     |                      |                    |                     |                   |                    |               |                    |                      |                      |              |
|  |                                 |   |                                      |                               |                               |                     |                               |                      |                             |                              |                     | _                   |                      | -                   |                    |                     |                      |                    |                     |                   |                    |               |                    | _                    |                      |              |
| RESULT 1 AA757305/c LOCUS DEFINITION ACCESSION VERSION KEYWORDS SOURCE ORGANISM  |                                 | 498<br>C 499<br>500   | 494<br>C 495<br>196                  | 491<br>492<br>493             | c 489                         |                     | C 484                         | 481<br>482<br>483    | c 479<br>c 480              |                              | C 475               |                     | 470<br>471           | c 468               | c 467              | C 464               | C 463                | C 461              | C 459               | C 457             | 455                | 453<br>454    | c 451<br>452       | C 449                | C 446                | 445          |
| E H H E A  |                                 |   | 17.6<br>17.6                         |                               | 17.6                          | 17.6<br>17.6        | 17.6<br>17.6                  | 17.6                 | 17.6<br>17.6                | 17.6                         | 17.6<br>17.6        | 17.6                | 17.6                 | 17.6<br>17.6        | 17.6<br>17.6       | 17.6                | 17.6                 | 17.6               | 17.6                | 17.6              | 17.6               | 17.6          | 17.6               | 17.6                 | 17.0                 | 17.          |
| A757305<br>A757305<br>AAGE:13269523 'S<br>F 12 (HUMAN);, mR<br>A757305<br>T.<br>T.<br>J. GI:280<br>J. GI |                                 | 0000  | 0000                                 | 000                           | 000                           | 000                 | 000                           | 000                  | 00                          | 000                          | 000                 | 000                 | 000                  | 00                  | 00                 | 00                  | 00                   |                    | 000                 | 000               | 000                | 000           | 0 0                | 000                  | 000                  | 0            |
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| Soares_NFL<br>52 3' simi<br>N);, mRNA<br>GI:280516   |                                 | 55 55 55 55<br>55 55 55 55<br>55 55 55 55<br>55 55                                  |                                      |                               |                               |                     |                               |                      |                             |                              |                     |                     |                      |                     |                    |                     |                      |                    |                     |                   |                    |               |                    |                      |                      |              |
| _ 1 ≯∃ਜ  |                                 | 9 9 9 9<br>A A A A  | 9 9 9 9 9 5<br>5 8 8 8 5             | 9 9 9<br>4 4 4                | 17<br>9 A                     | 9 A                 | 14                            | 14                   | 17                          | 999                          | 9 4 ×               | 17<br>9 A           | 17<br>17             | 10<br>9 A           | 9 9<br>A A         | 17<br>9 A           | 10<br>17             | 17                 | 17                  | 190               | o 00 0             | 999           | 17                 | 9 /                  |                      |              |
| 58<br>GBC<br>to<br>luenc   | ALIGNMENTS                      | A1/38849<br>AU263818<br>AA468168<br>AA611954  | 1251532<br>1267772<br>1340421        | A808779<br>A889400<br>T251270 | B04256<br>B700854             | 1251394<br>BQ273992 | BQ548919<br>AZ441898          | BM965665<br>BM965930 | B02174<br>B1491763          | 1567876                      | AZ435796<br>A645849 | AL761583<br>A790169 | AZ789268<br>AZ805887 | AV834440<br>A570437 | A914838<br>I799301 | A2876327<br>A668516 | AV948642<br>AZ817259 | BH864938           | AZ796868            | VU107918          | VII 1 0 6 0 1 A    | 10104970      | AZ804582           | A18651/3<br>AU060301 | AI023230<br>AI287232 |              |
| RNA<br>Sapien:<br>0_cds1   | <b>.</b> ,                      |   |                                      |                               |                               |                     |                               |                      |                             |                              |                     |                     |                      |                     |                    |                     |                      |                    |                     |                   |                    |               |                    |                      |                      |              |
| linear EST<br>ens cDNA clone<br> sl ZINC FINGER  |                                 | AI<br>AA<br>AA  | AI AI                                | , A A                         | A E -                         | . E                 | ~                             |                      |                             | A A A                        | .⊈.                 | ₽                   | ~ .                  | ⊉.                  | PP                 | ⊉.                  |                      |                    |                     | 2 2               | ב עי               | י סי          | >                  | <b>&gt;</b> >        | <b>&gt;</b> >        |              |
| EST<br>Lone<br>NGER  |                                 | 738849<br>1263818<br>1468168<br>611954  | 251532                               | 1808779<br>889400             | 700854                        | [251394<br>)Q27399  | 3054891<br>244189             | 3M96566              | 302174<br>3149176           | 1146706                      | 1645849             | AL76158<br>1790169  | AZ7892(              | 4V83444             | 4914831<br>1799301 | AZ8763:<br>A668516  | AV9486               | AZ9892<br>BH8649   | AZ4238<br>AZ7968    | 010791            | U10600             | U10497        | AZ8045             | 186517<br>U06030     | I02323<br>I28723     | вн8470       |
| ט ט  |                                 | AU26<br>nc731<br>vo86   | qv28<br>qv28<br>ap64<br>tb40         | nv74<br>am27                  | 38 2M0<br>CSRL-<br>zj36       | qv38                | 19 ik9                        | 55 ko1<br>601        | csRL-                       | 2 gw20<br>5 zo34             | 96 1M0              | 83 Ara<br>9 vv82    | 68 2MC               | 40 AV8              | 8 vy94<br>1 tw78   | 27 2MC<br>5 ac49    | 42 AVS<br>59 2M0     | 38 2M(             | 68 2MC              | # AU10            | 6 AU10             | 0 AU10        | 82 2MC             | 1 AU00               | 0 ov71               | 74 SAI       |
| 31-DEC-1998<br>ROTEIN  |                                 | AI738849 tr29g04.x<br>AU263818 AU263818<br>AA468168 nc73b11.r<br>AA611954 vo86c12.r | b02.x<br>h02.x<br>=12.x              | b07.r<br>f01.s                | 013M17<br>29f11-<br>al2.s     | b09.х<br>4с05.у     | ъсоз.у<br>3е02.у<br>234м15    | 5c09.y<br>9d01.y     | 149C12<br>4g01.w            | 612.s                        | 1223102<br>d02.r    | bidops<br>c09.r     | 036G20<br>067H03     | 34440<br>f08.s      | g10.r              | 191H14<br>d03.s     | )48642<br> 086I06    | )272H01<br>.K_0971 | )203P22<br>)052J14  | 7918              | 9009               | )4970         | )065009            | 9b01.x<br>50301      | 3d11.s               | K 0130       |

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                                 Emmert-Buck, M.D., Ph.D.

CDNA Library Preparation: Stratagene, Inc.

CDNA Library Arrayed by: Greg Lennon, Ph.D.

DNA Sequencing by: Washington University Genome Sequencing Center

DNA Sequencing by: Washington University Genome Sequencing Center

Clone distribution: NCI-GAP clone distribution information can be
                                                                                                                                                                                                                                                                                        Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalla; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 60) NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap. National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   of14e11.s1 NCI_CGAP_Kid6 Homo similar to gb:X16282_cds1 ZINC
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Seq primer: -40ml3 fwd. ET from Amersham
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                                                                                                                                                                                                                     Unpublished (1997)
Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AA834171.1 GI:2907770
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              mRNA sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
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                                                                                                                                                                    Tissue Procurement: L. Jeffrey Medeiros,
                                                                                                                                                                                                                                                                      Tumor Gene Index
                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens
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/ww-bio.llnl.gov/bbrp/image/image.html
                       found through the I.M.A.G.E. Consortium/LLNL at:
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(bases 1 to 58)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /note-"Organ: pooled; Vector: pT7T3D-Pac (Pharmacia) with a modified polylinker; Site_1: Not I; Site_2: Eco RI; a modified polylinker; Site_1: Not I; Site_2: Eco RI; Equal amounts of plasmid DNA from three normalized libraries (fetal lung NbHL19W, testis NHT, and B-cell NCI_CGAP_GCB1) were mixed, and ss circles were made in vitro. Following HAP purification, this DNA was used as tracer in a subtractive hybridization reaction. The driver was PCR-amplified cDNAs from pools of 5,000 clones made
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   from the same 3 libraries. The pools consisted of I.M.A.G.E. clones 297480-302087, 682622-687239, 726408-728711, and 729096-73139. Subtraction by I.M.A.G.E. charting Ronaldo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Soares and M. Fatima Bonaldo.
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/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /clone="IMAGE:1326952"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /organism="Homo sapiens"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   60 bp mRNA linear EST 09-JUN-1998 domo sapiens cDNA clone IMAGE:1421132 ZINC FINGER PROTEIN CLONE 647 (HUMAN);,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   59
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AA864103 58 bp mRNA linear EST 11-MAR-199 wn42e02.rl Stratagene mouse skin (#937313) Mus musculus cDNA clone IMAGE:1023866 5' similar to gb:D10632 Mouse mRNA for zinc finger
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Insert Length: 2306 Std Error: Seq primer: -40ml3 fwd. ET from High quality sequence stop: 1.
                                                                                                                                                                                            Contact: Marra M/Mouse EST Project WashU-HHMI Mouse EST Project
                                                                                                                                                                                                                                                                                                   Marra,M., Hillier,L., Allen,M., Bowles,M., Dietrich,N., Du
Geisel,S., Kucaba,T., Lacy,M., Le,W., Martin,J., Morris,M.
Schellenberg,K., Steptoe,M., Tan,F., Underwood,K., Moore,E
Theising,B., Wylie,T., Lennon,G., Soares,B., Wilson,R. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             protein (MOUSE);,
AA864103
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Emmert-Buck, M.D., Ph.D.
                                    This clone is available royalty-free through LLNL ; IMAGE Consortium (info@image.llnl.gov) for further i
                                                                                                      Tel: 314 286 1800
Fax: 314 286 1810
                                                                                                                                                 Washington University School of MedicineP
4444 Forest Park Parkway, Box 8501, St. Lo
                                                                                                                                                                                                                                                            The WashU-HHMI Mouse EST Project
                                                                                                                                                                                                                                                                                                                                                                                                              Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      EST
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                    MGI:574642
                                                                                                                                                                                                                                        Unpublished (1996)
                                                                                                                                                                                                                                                                                   Waterston, R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mus musculus
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                                                                                    .mouseest@watson.wustl.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /tissue_type="kidney tumor"
/lab_host="SOLR (kanamycin resistant)"
/note="Organ: kidney; Vector: Bluescript SK-; Site_1:
EcoRI; Site_2: XhoI; Cloned unidirectionally. Primer:
Oligo dT. Pooled kidney tumors. 5' adaptor sequence: 5'
GAATTCGGCACGAG 3' 3' adaptor sequence: 5'
CTCGAGTTTTTTTTTTTTTTTTTT 3' Average insert size: 1.0 kl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /clone="IMAGE:1421132"
/clone_lib="NCI_CGAP_Kid6"
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/db_xref="taxon:9606"
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80.0%;
-28ml3 revl ET from Amersham
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Pred. No.
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                                                             contact the
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; Murinae; Mus
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AA425976
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JOURNAL
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Kucaba, T., Allen, M., Bowles, L., Dubuque, T., Geisel, G., Jost, S., Schellenberg, K., Le, N., Lennon, G., Marra, M., Martin, J., Moore, B., T., Waterston, R. and Wilson, R. Washu-Merck EST Project 1997
Unpublished (1997)
Contact: 1977
                                                                                                                                                                                                                                            This clone is available royalty-free through LINL; contact the IMAGE Consortium (info@image.llnl.gov) for further information. Trace considered overall poor quality Possible reversed clone: similarity on wrong strand Seq primer: -28ml3 rev2 ET from Amersham High quality sequence stop: 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                 Washington University School of Medicine
4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                s
                                                                                                                                                                                                                                                                                                                                                                                                Tel: 314 286 1800
Fax: 314 286 1810
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Contact: Wilson RK
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        zw17b06.rl Soares ovary IMAGE:769523 5' similar
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Location/Qualifiers
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/tissue_type="ovarian tumor"
/tissue_type="ovarian tumor"
/lab_host="DH10B (ampicillin resistant)"
/note="Organ: ovary; Vector: pT7T3D (Pha
                                                                                                                  /organism="Homo sapiens"
/db_xref="GDB:5979393"
/db_xref="taxon:9606"
/clone="IMAGE:769523"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /lab_host="SOLR (kanamycin resistant)"
/note="organ: skin; Vector: pBluescript SK-; Site_1: EcoRI
/ Site_2: XhoI; Cloned unidirectionally, Primer: Oligo
dT. Whole skin from 11 week old C57BL/6 female mice.
Average insert size: 1.0 kb; Uni-ZAP XR Vector; -5'
adaptor sequence: 5' GAGTTTTTTTTTTTTTTTTTT 3'"
sequence: 5' CTCGAGTTTTTTTTTTTTTTTTTTTTT 3'"
                                                                             /sex="Female"
                                                                                          /clone_lib="Soares ovary tumor NbHOT"
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/dev_stage="11 weeks old"
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/clone="IMAGE:1023866"
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/strain="C57BL/6"
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15 g 14 t
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Pred. No. 1
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Nor NoHOT I
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L.9e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         www-bio.llnl.gov/bbrp/image/image.html
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CDNA Library Preparation: M. Bento Soares, Ph.D. CDNA Library Arrayed by: Greg Lennon, Ph.D. DNA Sequencing by: Washington University Genome Sequencing C Clone distribution: NCI-CGAP clone distribution information found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Emmert-Buck, M.D., Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Tissue Procurement: Christopher Moskaluk, M.D., Ph.D.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Tumor Gene Index
Unpublished (1997)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anat
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36; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Contact: Robert Strausberg, Ph.D.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 cgapbs-r@mail.nih.gov
                                                                                                                                                                                                     /clone="IMAGE:1632014"
/clone_lib="NCI_CGAP_Br2"
/sex="female, pooled"
/tissue_type="breast"
/lab_bost="DH10B"
                            /note="Vector: pT7T3D-Pac (Pharmacia) with a modified /note="Vector: pT7T3D-Pac (Pharmacia) with a modified polylinker; 1st strand cDNA was prepared from pooled bulk breast tumor tissue, and was then primed with a Not I - oligo(dT) primer. Double-stranded cDNA was ligated to Eco RI adaptors (Pharmacia), digested with Not I and cloned into the Not I and Eco RI sites of the modified pT7T3 vector. This library is the normalized version of NCI_CGAP_Br1.1. Library was constructed by Bento Soares and M. Fatima Bonaldo. "
                                                                                                                                                                                                                                                                                                            /organism="Homo sapiens"
/db_xref="taxon:9606"
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73.5%;
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Query Match

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Best Local S
Matches 38
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   57
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            vi54d09.x1 Beddington mouse embryonic region Mus musculus cDNA clone IMAGE:907601 3' similar to gb:x74855 m.musculus Zfp-51 mRNA for a zinc finger protein (MOUSE);, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Marra,M., Hillier,L., Kucaba,T., Martin,J., Beck,C., Wylie,T., Underwood,K., Steptoe,M., Theising,B., Allen,M., Bowers,Y., Person,B., Swaller,T., Gibbons,M., Pape,D., Harvey,N., Schurk,R., Ritter,E., Kohn,S., Shin,T., Jackson,Y., Cardenas,M., McCann,R., Waterston,R. and Wilson,R.

The WashU-NCI Mouse EST Project 1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity 39; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Washington University School of Medicine 4444 Forest Park Parkway, Box 8501, St. I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               This clone was previously sequenced on the 5' data is from the 3' end
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Unpublished (1999)
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mus musculus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Possible reversed clone: polyT not found
Seq primer: -28ml3 rev1 ET from Amersham
High quality sequence stop: 1
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       This clone is available royalty-free through LLNL; IMAGE Consortium (info@image.llnl.gov) for further i
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Contact: Marra M/WashU-NCI Mouse EST Project 1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MGI:528265
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                                                                                             Similarity

    (bases 1 to 58)

                                   CAGAGGACACATTCAGGGGAAAAAGCCTTTTATGTGCAGGGAGTGTGGCAGAAGGTTT 1802
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CAGAGAATTCATTCTGGAGAAAACCCTTCCAAAGTAAGTGAATGTGGCAAATCTTTT 1
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314 286 1810
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             mouseest@watson.wustl.edu
                                                                                                                                                                 from C57BL6 x DBA matings, excluding embryos that had developed head folds and all extraembryonic tissues. Average insert size: 1.3 kb (range: 0.5 - 3.0 kb). Referenced in Development 121, 2479-2489 (1995)"
                                                                                                                                                                                                                                                               /note="Organ: whole embryo; Vector: pCMV-SPORT; Site_1: Sall; Site_2: NotI; Cloned unidirectionally. Primer: Oligo dT. Gastrulating embryos were collected at 7.5dpc
                                                                                                                                                                                                                                                                                                                          /tissue_type="embryo"
/dev_stage="7.5dpc"
/lab_host="DH12S"
                                                                                                                                                                                                                                                                                                                                                                                                                        /db_xref="taxon:10090"
/clone="IMAGE:907601"
                                                                                                                                                                                                                                                                                                                                                                                                                                                            /organism="Mus musculus"
/strain="C57BL/6 x DBA"
                                                                                                                                                                                                                                                                                                                                                                                                  /clone_lib="Beddington
                                                                                                                                                                                                                                                                                                                                                                                  'sex="pooled"
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No. 6.7e+03;
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                                                                                                                                                                              AA136306 55 bp mrNA linear EST 30-NOV-1996
ZN89602.sl Stratagene lung carcinoma 937218 Homo sapiens cDNA clone
IMAGE:565346 3' similar to SW:KR4_HUMAN P10075 HKR4 PROTEIN;, mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   High quality sequence stop: 1
Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Christopher Moskaluk, M.D.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap
National Cancer Institute, Cancer Genome Ana
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
                                                                                               EST
                                                                                                                                            sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Contact: Robert Strausberg, Ph.D.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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                                              Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               www-bio.llnl.gov/bbrp/image/image.html
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               M. Fatima Bonaldo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /clone="IMAGE:1501686"
/clone_lib="NCI_CGAP_Kid3"
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/db_xref="taxon:9606"
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                                                                                                                   GI:1697533
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                        Euteleostomi;
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AZ513479/c
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Best Local Similarity
Matches 33; Conser
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1050 TGGAAGTCAAACCTGATCACACACATCAGAGGACACACTCAGGGGAGAAAACCTTAATG 1104
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           55
                                   Unpublished (2000)
Contact: Robert B.
                                                                                                                       Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., D. Islam, H., Longacre, S., Mahmoud, M., Meenen, E., M., Rose, M., Rose, R., Stokes, R., Tingey, A.,
                                                                                                                                                                                                   Eukaryota; Metazoa; Chordata; Mammalia; Eutheria; Rodentia;
                                                                                                                                                                                                                                                                                                                      1M0359P21R Mouse 10kb | clone UUGC1M0359P21 R, AZ513479
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 (bases 1 to 55)
Hillier, L., Lennon, G., Becker, M., Bonaldo, M.F., Chiapelli, B., Chissoe, S., Dietrich, N., DuBuque, T., Favello, A., Gish, W., Hawkins, M., Hulman, M., Kucaba, T., Lacy, M., Le, M., Mardis, E., Moore, B., Morris, M., Parsons, J., Prange, C., Rifkin, L., Rohlfing, T., Schellenberg, K., Soares, M.B., Tan, F., Thierry-Meg, J., Trevaskis, E., Underwood, K., Wohldmann, P., Waterston, R., Wilson, R. and Marra, M. Generation and analysis of 280,000 human expressed sequence tags
University of Utah
                                                                        plasmid inserts
                                                                                       Mouse whole genome
                                                                                                           and Wright, D., Weiss, R.
                                                                                                                                                                                   Mammalia; Eutheria;
1 (bases 1 to 45)
                                                                                                                                                                                                                                              Mus musculus
                                                                                                                                                                                                                                                                                                     AZ513479.1 GI:10694795
               University of Utah Genome Center
                                                                                                                                                                                                                                                              house mouse.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further information. Possible reversed clone: similarity on wrong strand Seq primer: -40M13 fwd. from Amersham
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                High quality sequence stop:
Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Contact: Wilson RK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Genome Res. 6 (9), 97044478
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Email: est@watson.wustl.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TGGCCTNCTCATCTNGNNNAGCATNAGAGNNTTCACTCAGGGGANAAGCCNTATG
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/tissue_type="lung carcinoma"
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/db_xref="GDB:4594411"
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                                                                                scaffolding with paired end reads from 10kb
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Pred.
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No. 1.1e+04;
                                                                                                                                                                                               Craniata; Vertebrata; Euteleostomi;
Sciurognathi; Muridae; Murinae; Mus
                                                                                                                     Beacorn,T., Duval,B., Hamil,C
,M., Meenen,E., Pedersen,T., Ro
R., Tingey,A., von Niederhause
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                carcinoma 937218"
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VERSION
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AA948106/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1419 CACACAGGAGAGAGCCTTATGTATGCACAGAATGTGGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       44 CATACCGGGGAGAAACCCTATCTATGTAAACAATGTGGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           30;
                Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Christopher Moskaluk, M.D.,
Emmert-Buck, M.D., Ph.D.
                                                                                              Tumor Gene Index
Unpublished (1997)
                                                                                                                                                                                                                                                                                                                                                                 on5la04.sl
similar to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Rm. 308, Biomedica
84112, USA
Tel: 801 585 5606
Fax: 801 585 7177
                                                                             Contact: Robert Strausberg, Ph.D.
                                                                                                                                                          NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap
                                                                                                                                                                                                                       Eukaryota; Metazoa;
                                                                                                                                                                                                                                            Homo sapiens
                                                                                                                                                                                                                                                                                      EST
                                                                                                                                                                                                                                                                                                       AA948106.1 GI:3109359
                                                                                                                                                                                                                                                                                                                              AA948106
                                                                                                                                                                                                                                                                                                                                                                                                         AA948106
                                                                                                                                        National Cancer Institute, Cancer
                                                                                                                                                                                                      Mammalia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Erro
Plate: 0359 row: P column: 2)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity
cDNA Library Preparation: M. Bento Soares, Ph.D
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         High quality sequence stop: 45.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Class: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Seq primer: CACACAGGAAACAGCTATGACC
                                                                                                                                                                                 (bases 1 to 50)
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USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from derivative of pWD42 (gil4732114) gbjn47129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     was hydrodynamically sheared by repeated passage through 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with 74 DNA polymerase and 74 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"/note="Vector: PW042nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Biomedical Polymers Research Bldg.,
                                                                                                                                                                                                      Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (http://www.jax.org/resources/documents/dnares/).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /sex="Male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /clone_lib="Mouse 10kb plasmid UUGC1M library"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /db_xref="taxon:10090"
/clone="UUGC1M0359P21"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /organism="Mus musculus"
/strain="C57BL/6J"
                                                                                                                                                                                                                                                                                                                                                         50 bp mRNA linear EST 23 NCI_CGAP_CO8 HOMO Sapiens cDNA clone IMAGE:1560 gb:M58297 ZINC FINGER PROTEIN 42 (HUMAN);, mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0.98;
                                                                                                                                                                                                  Chordata; Craniata; Vertebrata; Euteleostomi; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 24.6; DB
Pred. No. 2.3e+(
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                                   Ph.D.,
                                                                                                                                                                                                                                                                                                                                                                              ar EST 23-JUN-1998
IMAGE:1560174 3'
                                                                                                                                        Project (CGAP),
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R37288/c
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VERSION
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             50
                                                                                                                                                                                                                                                                                                                                                                               ,M., Hultman,M., Kucaba,T., Le,M., Lenmon,G., Marra,M.; Parsons,J. Rifkin,L., Rohlfing,T., Soares,M., Tan,F., Trevaskis,E., Waterston,R., Williamson,A., Wohldmann,P. and Wilson,R.

The WashU-Merck EST Project
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mammalia; Eutheria; Primates; Catarrhini; Homi (bases 1 to 38)
Hillier, L., Clark, N., Dubuque, T., Elliston, K.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        R37288 38 bp mRNA yf67a06.sl Soares infant brain 1NIB Homo IMAGE:27044 3' similar to gb:X16282_cds1
                                                           Insert Size: 690
High quality sequence starts: 1 High quality sequence stops: 1
Source: IMAGE Consortium, LLNL This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further information. Trace considered overall poor quality Insert Length: 690 Std Error: 0.00
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                                                                                                                                                                                                                                                                                  Washington University School of Medicine 4444 Forest Park Parkway, Box 8501, St. 1
                                                                                                                                                                                                                                                                                                                                                             Unpublished (1995)
                                                                                                                                                                                                                                     Tel: 314 286 1800
Fax: 314 286 1810
                                                                                                                                                                                                                                                                                                                                    Contact: Wilson RK
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                                               Seq primer: -21m13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        human
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GCACCAGCGCATCCACCACGGCGAGAAGCCCTTCGCGTGCCCCGAGTGCG
                      quality sequence stop: 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                          est@watson.wustl.edu
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     modified polylinker; 1st strand cDNA was prepared from colon adenocarcinoma, and was then primed with a Not I - oligo(dT) primer. Double-stranded cDNA was ligated to Eco RI adaptors (Pharmacia), digested with Not I and cloned into the Not I and Eco RI sites of the modified pT7T3 vector. Library is normalized. Library was constructed by Bento Soares and M. Fatima Bonaldo. "
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /clone="IMAGE:1560174"
/clone_lib="NCI_CGAP_CO8"
/tissue_type="adenocarcinoma"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /organism="Homo sapiens"
/db_xref="taxon:9606"
Location/Qualifiers
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Pred. No. 2.8e+04;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     36 AAGCCCTATGAGTGTAATGANTGTGGGAAGGCCTTT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           28;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
Plate: LLAM11857 row: a column: 02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Lothar Hennighausen Ph.D., Chu-Xia Den
cDNA Library Preparation: Life Technologies, Inc.
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    603296901F1 NCI_CGAP_Mam3 Mus
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Contact: Robert Strausberg, Ph.D.
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
1. (bases 1 to 59)
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Library constructed by Life Technologies. Investigators providing samples: Lothar Hennighausen/Chu-Xia Deng, NIH Reference for transgenic model: Xu et al., Nature Construction and Construction 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        constructed by Bento Soares and M.Fatima Bonaldo 12.c 5 g 10 t 1 others
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/db_xref="GDB:399391"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                               /clone_lib="NCI_CGAP_Mam3"
/tissue_type="tumor, gross tissue"
/dev_stage="10 months"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /organism="Mus musculus"
/strain="129,C57BL/6J,FVB/N"
/db_xref="taxon:10090"
/clone="IMAGE:5337481"
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/note="Organ: whole brain; Vector: Lafmid BA;
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Pred. No. 3.5e+04;
0; Mismatches 8;
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s musculus cDNA clone IMAGE:5337481 5',
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AA867748
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                  Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            54 GCTGCTTTCAGCCCGCCTCAAGACTGACAGCCTGCGCAGTTCA 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 (bases 1 to 46)
Marra, M., Hillier, L., Allen, M., Bowles, M., Dietrich, N., Dub
Geisel, S., Kucaba, T., Lacy, M., Le, M., Martin, J., Morris, M.,
Schellenberg, K., Steptoe, M., Tan, F., Underwood, K., Moore, B.
Theising, B., Wylie, T., Lennon, G., Soares, B., Wilson, R. and
                                                                                                                                                                                                                                                                                                                                                                                   Trace considered overall poor quality Possible reversed clone: similarity on wrong strand Seq primer: -28ml3 rev2 ET from Amersham High quality sequence stop: 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Washington University School of MedicineP
4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AA867748

46 bp mRNA linear EST 16-M/VX16a12.rl Soares_thymus_2NbMT Mus musculus cDNA clone IMAGE:1264606 5' similar to SW:Z154_HUMAN Q13106 ZINC FINGER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Unpublished (1996)
                                                                                                                                                                                                                                                                                                                                                                                                                                               MGI:667158
                                                                                                                                                                                                                                                                                                                                                                                                                                                       This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further information.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Tel: 314 286 1800
Fax: 314 286 1810
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Contact: Marra M/Mouse EST Project
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The WashU-HHMI Mouse EST Project
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Waterston, R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mus musculus
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                                                      Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       mouseest@watson.wustl.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ۵
                                                                                                                                                                                                                                                                                             /organism="Mus musculus"
/strain="C57BL/6J"
/db_xref="taxon:10090"
/clone="IMAGE:1264606"
                                                                                                                                                                                                                       /dev_stage="4 weeks"
/lab_host="DH10B"
                                                                                                                                                                                                                                                    /tissue_type="Thymus"
                                                                                                                                                                                                                                                                     /sex="male"
                                                                                                                                                                                                                                                                              /clone_lib="Soares_thymus_2NbMT"
                                                                                                                                                                                                                                                                                                                                                                      Location/Qualifiers
69.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0.98;
72.18;
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   . 69;
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Pred. No. 4.7e+04;
0; Mismatches 12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  23
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 23.6;
No. 4
 .6; DB 9;
. 4.6e+04;
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               Length 46;
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                                                      RESULT 15
BF638864
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                                         DEFINITION
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                                                                                                                                                                                       Local Similarity
les 33; Conserv
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ACCESSION
VERSION
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                                                                                                                                                                                                                  1052 GAAGTCAAACCTGATCACACATCAGAGGACACACTCAGGGGAGAAACCT 1100
                      60 bp
NF060H02PL1F1026 Phosphate starved
clone NF060H02PL 5', mRNA sequence.
BF638864
BF638864
BF638864.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Washington University School of Medicine 4444 Forest Park Parkway, Box 8501, St. Louis, MO Tel: 314 286 1800
Fax: 314 286 1810
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AA227017

49 bp mRNA linear EST 11-MAR-1: zr19c09 rl Stratagene NT2 neuronal precursor 937230 Homo sapiens cDNA clone IMAGE:663856 5' similar to SW:ZN31_HUMAN P17040 ZINC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          32;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Contact: Wilson RK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      White,Y., Wylie,T., Watersto
WashU-NCI human EST Project
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Krizman,D., Kucaba,T., Lacy,M.,
,J., Moore,B., Schellenberg,K.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Hillier,L., Allen,M., Bowles,L.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Unpublished (1997)
                                                                                                                                                                             GAATAGAAGTCTTATTGAACATCAGAAAATCCACACTGGTGAGAAACCT 49
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AGGGGAAAAGCCTTTTATGTGCAGGGAGTGTGGCAGAAGGTTTCGG 1805
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 primer: -28m13 rev1 ET from Amersham
quality sequence stop: 40.
                                                                                                                                                                                                                                                             Conservative
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                                                                                                                                                                                                                                                                                                                                                        /clone_lib="Stratagene NT2 neuronal precursor 937230"
/tissue_type="neuroepithelial cells"
/dev_stage="Ntera-2 neuroepithelial cells"
/lab_host="SOLR (kanamycin resistant)"
/note="Organ: brain; Vector: pBluescript SK-; Site_1:
ECORI; Site_2: Xho; Cloned unidirectionally. Primer:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /organism="Homo sapiens"
/db_xref="GDB:5589788"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /clone="IMAGE:663856"
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                                                                                                                                                                                                                                                                             0.8%;
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                                                                                                                                                                                                                                                             0;
                                                                                                                                                                                                                                                         Score 23.4; DB 9;
Pred. No. 5.5e+04;
0; Mismatches 16
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                                                       mRNA
leaf
                                                                                                                                                                                                                                                           16;
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                                                       Medicago truncatula
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Email: mjharrison@noble.org
Insert Length: 60 Std Error: 0.00
Plate: 060 row: H column: 02
Seq primer: TCACACAGGAAACAGCTATGAC.
Location/Qualifiers
Contact: Wilson RK
Washington University School of Medicine
4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108
                                                                                        Hillier, L., Clark, N., Dubuque, T., Elliston, K., Hawkins, M., Holman, M., Kucaba, T., Le, M., Lennon, G., Marra, M., Parsons, J. Rifkin, L., Rohlfing, T., Soares, M., Tan, F., Trevaskis, E., Waterstor, R., Williamson, A., Wöhldmann, P. and Wilson, R.
                                                                                                                                                                             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalla; Eutherla; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 37)

Hillier,L., Clark,N., Dubuque,T., Elliston,K., Hawkins,M., Holman
                                                                                                                                                                                                                                                                                                                                                                                             yd39a12.rl Soares fetal
IMAGE:110590 5' similar
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      l Similarity
30; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2510 Sam Noble Parkway, Ardmore,
Tel: 580 221 7325
Fax: 580 221 7380
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The Samuel Roberts Noble Foundation

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Medicago truncatula
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots;
Rosidae; eurosids I; Fabales; Fabaceae; Papilionoideae; Trifolieae;
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barrel medic.
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                                                                         Unpublished (1995)
                                                                                                                                                                                                                                                                 Homo sapiens
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17 c 12 g 9 t 7 others
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /clone_lib="Phosphate starved leaf"
/tissue_type="leaf"
/dev_stage="trifoliate"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /organism="Medicago truncatula"
/db_xref="taxon:3880"
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ver spleen lNFLS Homo sapiens cDNA clone
gb:X52354 ZINC FINGER ΡΡΟΜΠΤΥ:
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Email: est@watson.wustl.edu
Insert Size: 752
High quality sequence stops: 1
High quality sequence stops: 1
Source: IMAGE Consortium, LLNL This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov)
--- forther information. Trace considered overall poor quality
--- forther information: 314 Error: 0.00
riale: 0091 row: I column: 11
Seq primer: CGTTGTAAAACGACGGCCAGT
Class: plasmid ends
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27; Conserv
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Location/Qualifiers
                                                                                                                                          University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Res
                                                                                                                                                                                                                                                               Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly
,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A.
and Wright,D., Weiss,R.
                                                                                                                                                                                                                                                                                                                  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus 1 (bases 1 to 54)

Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1M0091111F Mouse 10kb plasmid UUGC1M library Mus musculus
                                                                                         Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                               Contact: Robert B.
                                                                                                                                                                                                                                                                                                                                                                                           Mus musculus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               clone UUGC1M0091111 F, DNA sequence.
                                                      Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Erro
                                                                                                                                                                                                                Unpublished (2000)
                                                                                                                                                                                                                                plasmid inserts
                                                                                                                                                                                                                                                 Mouse whole genome scaffolding with paired end reads from 10kb
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/clone="IMAGE:110590"
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73.0%;
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Pred. No. 6.9e+04;
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National Cancer Institute, Cancer Genome Anatomy
Tumor Gene Index
                  Trace considered overall poor quality Insert Length: 1396 Std Error: 0.00
                                                                        www-bio.llnl.gov/bbrp/image/image.html
                                                                                    cDNA Library Preparation: Life Technologies, Inc. cDNA Library Arrayed by: Greg Lennon, Ph.D. DNA Sequencing by: Washington University Genome Sequencing Center Clone distribution: NCI-CGAP clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                          Tissue Procurement: Christopher Moskaluk, M.D., Ph.D., Emmert-Buck, M.D., Ph.D.
                                                                                                                                                                                                                                Email: cgapbs-remail.nih.gov
                                                                                                                                                                                                                                                    Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                Unpublished (1997)
                                                                                                                                                                                                                                                                                                                                                                Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; I
Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AI687537 55 bp mRNA linear EST 14-DEC::999 tp88f10.x1 NCI_CGAP_Ut3 Homo sapiens cDNA clone IMAGE:2206411 3 similar to TR:O43516 043516 WASP INTERACTING PROTEIN. ; , mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              sequence. AI687537
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        primer:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (http://www.jax.org/resources/documents/dnares/). The INA
was hydrodynamically sheared by repeated passage through a
0.005 inch orifice at constant velocity. The sheared DNA
was blunt end-repaired with T4 DNA polymerase and T4
polynucleotide kinase. Adaptor oligonucleotides were
ligated to the blunt ends in high molar excess. The
adaptored DNA was purified and size-selected for a 9.5 to
10.5 kb range using preparative agarose gel
electrophoresis. Vector DNA was prepared from a derivative
of pWD42 (gi14732114|gbhAF129072.1), a copy-number
inducible derivative of plasmid R1. The vector was ligated
with adaptors complementary to the insert adaptors and
purified. The sheared, adaptored mouse DNA was annealed to
chemically-competent E. coli xL10-Gold (Stratagene) cells
and selected for ampicillin resistance."
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /lab_host="E. Coli strain XL10-Gold,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /clone_lib="Mouse 10kb plasmid UUGC1M library"
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/clone="UUGC1M0091111"
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/strain="C57BL/6J"
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Pred. No. 9e+04;
0; Mismatches 17;
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                                                                                                                                                                                  This sequence is a single read and was generated as scale clone-end sequencing project of the Tetraodon genome. For more information, please take a look at http://www.genoscope.cns.fr/Tetraodon.
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                                                                                                                                                                                                                                                                           Direct Submission
Submitted (12-APR-2000)
                                                                                                                                                                                                                                                                                                                          Genoscope.
                                                                                                                                                                                                                                                                                                                                                                  Unpublished
                                                                                                                                                                                                                                                                                                                                                                                    Charaterization and repeat analysis of the compact genome of the freshwater pufferfish Tetraodon nigroviridis
                                                                                                                                                                                                                                                                                                                                                                                                                               Weissenbach,
                                                                                                                                                                                                                                                                                                                                                                                                                                     Roest-Crollius,H., Jaillon,O., Dasilva,C., Fizames,C., Fisher,C., Bouneau,L., Billault,A., Quetier,F., Saurin,W., Bernot,A. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human gene number estimate provided by genome wide analysis using Tetraodon nigroviridis DNA sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 (bases 1 to 56)
Roest-Crollius, H., Jaillon, O., Dasilva, C., Boung
Bernot, A., Fizames, C., Wincker, P., Brottier, P.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Actinopterygii; Neopterygii; Teleostei; Buteleostei; Neoteleostei; Acanthomorpha; Acanthopterygii; Percomorpha; Tetraodontiformes;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GSS; genome survey sequence. Tetraodon nigroviridis. Tetraodon nigroviridis
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AL175368
AL175368.1 GI:7813425
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Unpublished
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Saurin, W. and Weissenbach, J
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Location/Qualifiers
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                    Ω
/Clone- ..../Clone-lib="G"
/Clone_lib="G"
/note="Genoscope sequence
/note="Genoscope sequence
                                                                              /organism="Tetraodon nigroviridis"
/db_xref="taxon:99883"
/clone="222K13"
                                                                                                                                                                location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /note="Organ: uterus; Vector: pCMV-SPORT6; Site_1: Sal1; Site_2: NotI; Cloned unidirectionally. Primer: Oligo dT. Average insert size 1.45 kb. Life Technologies catalog #: 11541-018"
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/db_xref="taxon:9606"
/clone="InAGE:2206411"
/clone_lib="NCI_CGAP_Ut3"
/tissue_type="poorly-differentiated endometrial adenocarcinoma, 2 pooled tumors"
/lab_host="DH10B"
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Pred. No. 9.1e+04;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Email: dmelton@biohp.harvard.edu
Library was constructed by Dr. Douglas Melton DNA sequencing by:
Washington University Genome Sequencing Center For information on
obtaining a clone please contact: Juliana Brown
(brown@fas.harvard.edu) This sequence now available from the IMAGE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Endocrine Pancreas Consortium
Harvard University, Howard Hughes Medical
Dept of Molecular and Cellular Biology, 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Unpublished (2000)
Other_ESTs: 1e65a07.y1
Contact: Douglas Melton, Klaus H. Kaestner, & Hiroshi
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              consortium, for clone orders contact: info@image.llnl.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Fax: 617-495-8557
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                                                                                  20. Single-stranueu .........by hydroxyapatite chromatography library."

1 5 c 12 g 19 t
                                                                                                                                                                                      Site_2: Sal 1; Starting library constructed using SuperScript Plasmid Library kit (Life Technologies).cDNA made by oligo-dr priming. Size-selected by column fractionation; average insert size 1.08 kb. Library was amplified once on solid support and plasmid DNA from library was prepared. The library DNA was normalized by method #4 from Bonaldo, Lennon, and Soares 1996 Genome Research 6:791-806; 0.5 microgram single-stranded library plasmid DNA was mixed with 5 micrograms PCR product representing library inserts and hybridized to an Ecot of 20. Single-stranded (unhybridized) plasmids were isolated by the december of the content of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /lab_host-"DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /tissue_type="Islets of Langerhans"
/dev_stage="Adult"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /sex="Both"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /organism="Homo sapiens"
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Pred. No. 9
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22.6; DB 13;
No. 1e+05;
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and used to make
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                       Length 55;
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                                                                                     oa44a03.sl t
similar to 7
AA767515
                                                     56 CAAAACACGGCCTACACACCCGGGGGAGAAGCCATACAAGTTTTCAAATTGTG 4
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                                                                                                                                                                                               Similarity
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2710 TCCCCTCTCCACTGAGAGCTGTTCTTTTGCTCAATAAAATTCTTT 2754
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            31;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Trace considered overall poor quality
Insert Length: 1430 Std Error: 0.00
Seq primer: -40ml3 fwd. ET from Amersham
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DNA Sequencing by: Washington University Genome Clone distribution: NCI-CGAP clone distribution found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Unpublished (1997)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AA767515.1 GI:2818530
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Bonaldo, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Tissue Procurement: Louis M. Staudt, M.D., Ph.D., Gerald Marti, M.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Tumor Gene Index
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap
National Cancer Institute, Cancer Genome Ana
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             www-bio.llnl.gov/bbrp/image/image.html
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cDNA Library Preparation: M. Bento Soares, Ph.D.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                cDNA Library Arrayed by: Greg Lennon, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (bases 1 to 58)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                          and Eco RI sites of the modified pT7T3 vector. L. went through one round of normalization, and was constructed by Bento Soares and M. Fatima Bonalds 13 c 15 g 20 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                /clone="IMAGE:1307788"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /organism="Homo sapiens"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                  /lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                    /tissue_type="germinal center B cell"
                                                                                                                                                                                                                                                                                                                                                                                                                          /clone_lib="NCI_CGAP_GCB1"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     sequence stop:
0.8%;
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22.6;
No. 1.
DB 9;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ph.D.,
                 Length 58;
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0; Pred.

Mismatches

Indels

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Gaps

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ACCESSION
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AA227334
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AUTHORS
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AL757225/c
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                                                                                                                                                                                                                                                                                                                                          Local
                                                                                                                                                                                                                                            2468 AACCTGTCTTCCCATTTGGTGTGCTTTCCTCCGATTG 2504
                                                                                                                                                                                                                            42
                zr17b11.rl Stratagene NT2 neuronal precursor 977230 Homo sapiens cDNA clone IMAGE:663645 5' similar to TR:G1049295 G1049295 KRAB ZINC FINGER PROTEIN ;, mRNA sequence.
                                                                                                                                                                                                                                                                                                                  l Similarity
28; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Submitted (17-JUN-2002) Weisshaar B., Max-Planck-Institut fuer Zuechtungsforschung, Carl-von-Linne-Weg 10, Koeln, 50829, Germany This sequence is recovered from the right border of the T-DNA. It indicates an insertion within the locus defined by clone F19K16. The sequences are generated at the MPI for Plant Breeding Research in the context of the GABI-Kat project. GABI-Kat is part of the
AA227334.1 GI:1848937
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Arabidopsis thaliana
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots;
Rosidae; eurosids II; Brassicales; Brassicaceae; Arabidopsis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        German Plant Genomics program designated 'GABI'. Information line availability can be found at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Strizhov, N., Rosso, M., Li, Y. and Weisshaar, B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Rosso,M., Strizhov,N., Li,Y., Reiss,B., Dekker,K. and Weisshaar,B. A new Arabidopsis thaliana T-DNA mutagenised population (GABI-Kat) for flanking sequence tag based reverse genetics
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Direct Submission
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Unpublished
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Unpublished
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Strizhov,N., Li,Y., Rosso,M., Viehoever,P., Dekker,K., Saedler,H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             genomic survey sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A pipeline for automated high-throughput generation of FSTs (flanking sequence tags) from Arabidopsis thaliana T-DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           thale cress.
                                                                                                                                                                                                                       AACATGTGATTGCATATGGTGTGTTTTCCTTTTATTG
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                                                                                                                                                                                                                                                                                                                  Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    insertions. The DNA fragment(s) resulting from the PCR were directly sequenced to determine the genomic sequence flanking the insertion. Sequences displaying significant similarity to the A. thalaina nuclear genome sequence were processed for submission. T-DNA derived sequences were
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /Clone_lib="Arabidopsis thaliana T-DNA insertion lines"
/note="PCR was performed on DNA from Arabidopsis thaliana
plants (T1) which were transformed with the T-DNA from
vector pAC161. The lines contain one or more T-DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /organism="Arabidopsis thaliana"
/strain="Columbia 0"
/db_xref="taxon:3702"
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                                                                                                                                                                                                                                                                                                                                     0.8%;
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.Pred. No. 1.1e+05;
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                                                                                                                                                                                                                                                                                                                                                        17;
                                                                                                                                                                                                                                                                                                                                                      Length 60;
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                                                                                                       EST 11-MAR-1998
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AUTHORS
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                                                                                                                                                                                                                                                                                                                                                                 RESULT 24
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ORGANISM
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JOURNAL
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         206
                                                                                                                                                                                                                                                                                 BM566527
kj63e09.yl Ascaris suum male gut
                                            McCarter, J., Clifton, S., Chiapelli, B., Pape, D., Martin, J., Wylie, T.,
Dante, M., Marra, M., Hillier, L., Kucaba, T., Theising, B., Bowers, Y.,
Gibbons, M., Ritter, E., Bennett, J., Franklin, C., Tsagareishvili, R.,
Ronko, I., Kennedy, S., Maguire, L., Beck, C., Underwood, K., Steptoe, M., Allen, M., Person, B., Swaller, T., Harvey, N., Schurk, R., Kohn, S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          30;
                                                                                                                                                                                                                                      EST.
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                    , Shin,T., Jackson,Y., Cardenas,M., McCann,R., Waterston,R. and
Wilson,R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Maumdial, Joseph (1998)

I (bases 1 to 43)

I (bases 1 to 43)

Rizman, D., Rucaba, T., Lacy, M., Lennon, G., Marra, M., Martin, J., Moore, B., Schellenberg, K., Steptoe, M., Tan, F., Theising, B., White, Y., Wylie, T., Waterston, R. and Wilson, R.
The Washington Univ. Nematode EST Project, 1999
                                                                                                                                                                           Eukaryota; Metazoa; Nematoda; Chromadorea; Ascaridida;
                                                                                                                                                                                               Ascaris suum
                                                                                                                                                                                                                    pig roundworm.
                                                                                                                                                                                                                                                     BM566527.1 GI:18827611
                                                                                                                                                                                                                                                                          BM566527
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further information. Possible reversed clone: similarity on wrong strand
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Washington University School of Medicine 4444 Forest Park Parkway, Box 8501, St.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Seq primer: -28m13 rev1 ET from Amersham
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Email: est@watson.wustl.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Contact: Wilson RK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Unpublished (1997)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Insert Length: 1843
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     human.
                                                                                                                                                                                                                                                                                                                                                                                                                                  GGCGTTCGTGGCATTCAGGGATGTGGCTGTGTGTACTTCACCCAG 248
                                                                                                                                                              Ascarididae; Ascaris.
                                                                                                                                                                                                                                                                                                                                                                                                                   GGATTCAGTGACGTTTGATGATGTGGCTGTGGAGTTCCCCCAG
                                                                                                                                           (bases 1 to 52)
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/db_xref="GDB:5589575"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /clone_lib="Stratagene NT2 neuronal precursor 937230"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /clone="IMAGE:663645"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /db_xref="taxon:9606"
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Pred.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
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No. 1.2e+05;
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                                                                                                                                                                                                                                                                                       mRNA sequence.
                                                                                                                                                                                                                                                                                                      SL1 TOPO v1 Murphy Chiapelli
                                                                                                                                                                                                                                                                                                                            mRNA
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REFERENCE
AUTHORS
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AZ310125/c
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Unpublished (1999)
Contact: McCarter JP
Contact: McCarter JP
The Washington Univ. Nematode EST Project, 1999
Washington University School of Medicine
4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108, USA
                                                                                                    Contact: Robert B. University of Utah University of Utah
                                                                                                                                                                                                                      Islam,H., Longacre,S., Mahmoud,M.,
M., Rose,M., Rose,R., Stokes,R.,
and Wright,D.,Weiss,R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                              AZ310125
1M0018M20R Mouse 10kb p
clone UUGC1M0018M20 R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The library was constructed by Claire Murphy, Brandi Chiapelli, Dr. James McCarter at Washington University, St. Louis. DNA Sequencing by: Washington University Genome Sequencing Center.
                                                                                                                                                                                   Mouse whole genome scaffolding with paired end reads from 10kb plasmid inserts
                                                                                                                                                                                                                                                                                                   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; 1 (bases 1 to 48)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Tel: 314 286 1800
Fax: 314 286 1810
                      Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                Unpublished (2000)
                                                                                                                                                                                                                                                                Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C.,
Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Rei
                                                                                                                                                                                                                                                                                                                                                                  Mus musculus
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                                                                                                                                                                                                                                                                                                                                                                                         house mouse
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ATCAACCCTGATCACGCACCAGAGGACACACTCAGGGGAGAAG 1685
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                                                                                     308,
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    ddunn@genetics.utah.edu
                                                                USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /note="Vector: pCRII-TOPO (Invitrogen): Site_1: EcoRI; Site_2: EcoRI; The library was constructed by Claire Murphy, Brandi Chiapelli, and Dr. James McCarter at Washington University, St. Louis. Oligo(dT)-SLI PCR based library. Ascaris suum male intestine cDNA PCR products of size >400 nucleotides containing SLI on the 5 end and oligo(dT) on the 3 end were non-directionally cloned into pCRII-TOPO(Invitrogen) following the Topo TA cloning
                                                                               Biomedical Polymers Research Bldg.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Baltimore, MD"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            protocol. Dissected nematode tissues were provided by Dr
Alan Scott (ascott@jhsph.edu) of the School of Public
Hygiene and Public Health at John Hopkins University in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /organism="Ascaris suum"
/db_xref="taxon:6253"
/clone_lib="Ascaris suum male
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Chiapelli McCarter"
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/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /sex="Male'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      location/Qualifiers
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                                                                                                                        Genome Center
                                                                                                                                         Weiss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0;
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Pred. No. 1.3e+05;
0; Mismatches 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  plasmid UUGC1M library Mus musculus genomic
, DNA sequence.
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                                                                                 2030 E., SLC,
                                                                                                                                                                                                                                                                                                                                           Euteleostomi;
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                                                                                                                                                                                                                                                                                                                          Murinae;
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AUTHORS
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ORGANISM
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AZ665324/c
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                                                                                                                                           COMMENT
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           46
                                                                                                                                                                                                                        Dunn, D., Aoyagi, A., Barber, M., Beucca., ...
Islam, H., Longacre, S., Mahmoud, M., Meenen, E.,
M., Rose, M., Rose, R., Stokes, R., Tingey, A., V
                                                                                                 University of Utah Genome University of Utah
                                                                                                                                                                                                                                                                                                                                                                                                         GSS
                                                                                                                                                                                                                                                                                                                                                                                                                        AZ665324
AZ665324.1 GI:11802470
                                                                                                                                                                                                                                                                                                                                                                                                                                                       AZ665324 55 bp DNA linear GSS 14-DEC-200
1M0546007R Mouse 10kb plasmid UUGC1M library Mus musculus genomic
clone UUGC1M0546007 R, DNA sequence.
                    Tel: 801 585 5606
Fax: 801 585 7177
                                                                                 Rm.
                                                                                                                                                              Unpublished (2000)
                                                                                                                                                                                 plasmid inserts
                                                                                                                                                                                                   Mouse whole genome scaffolding with paired
                                                                                                                                                                                                                                                                                                                  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae;
                                                                                                                                                                                                                                                                                                                                                              Mus musculus
                                                            84112; USA
                                                                                                                                         Contact: Robert B.
                                                                                                                                                                                                                                                                                                   Mammalia; Eutheria;
1 (bases 1 to 55)
                                                                                                                                                                                                                                                                                                                                                                                     house mouse
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GGGACTCGGCCAGTGGCTACAGGCCGCCCCCGCGGCCAACAGAGAGG
                                                                                 308,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
ddunn@genetics.utah.edu
                                                                                 Biomedical
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Mismatches

Indels

0;

Gaps

0,

Weiss

end

reads

from 10kb

Duval, B., Hamil, C

Euteleostomi;

Murinae;

Mus

GSS 14-DEC-2000

Pedersen, T.,

Reilly

Niederhausern, A.

Center

Polymers Research Bldg.,

20

s.

2030 E.,

SLC,

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Class: plasmid ends
High quality sequence stop: 48.
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                                                                                                                                                                                                                                            was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pND42 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored modes DNA was annealed to
                                                                                                                           chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance." 19 \ c \qquad 16 \ g \qquad 10 \ t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through 0.005 inch orifice at constant velocity. The sheared DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /lab_host="E. Coli strain XL10-Gold, T1-resistant, F
/note="Vector: PWD42nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jack
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /organism="Mus musculus
/strain="C57BL/6J"
/db_xref="taxon:10090"
                                                                                                                                                                                                                     adaptored vector DNA, and transformed into
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Laboratory Mouse DNA Resource
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /sex="Male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /clone="UUGC1M0018M20"
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   0.8%;
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Score 22; Pred. No.
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DB 17;
1.4e+05;
                             Length 48
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Jackson -
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FEATURES
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AI051772
                                                                                                                                                                                                                         COMMENT
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TITLE
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Best Local
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                           source
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    34;
                                                                     TMAGE Consortium (info@image.linl.gov) for further information. Trace considered overall poor quality Seq primer: -40ml3 fwd. Erf from Amersham High quality sequence stop: 1.
                                                                                                                                                                                         Unpublished (1997)
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ox45e08.sl Soares_total_fetus_wb2HF8_9w Homo sapiens CNNA clone IMAGE:1659302 3' similar to TR:065358 065358 HYPOTHETICAL PROTEIN contains MSR1.t2 TAR1 repetitive element; mRNA sequence AI051772
                                                                                                                                                              This clone is available royalty-free through LLNL;
                                                                                                                                                                                                                                                                               NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anat
                                                                                                                                                                                                                                                                                                                                                       Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AI051772.1 GI:3307306
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                                                                                                                                                                                                                                                                                                                                        (bases 1 to 58)
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                                                                                                                                                                                                                                                                   Gene Index
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."

13 c 14 g 15 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pND42 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid RI. The vector was ligated with adaptors complementary to the insert adaptors and
                                                 Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The
/organism="Homo sapiens"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (http://www.jax.org/resources/documents/dnares/). The DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /sex="Male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /clone_lib="Mouse 10kb plasmid UUGC1M library"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /clone="UUGC1M0546007"
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Pred. No. 1.6e
0; Mismatches
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olumn: 07
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. 1.6e+05;
tches 20;
                                                                                                                                                                                                                                                                                  Genome Anatomy Project (CGAP),
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FEATURES
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Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AUTHORS
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                                                                                                                                                                                        source
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             High quality sequence stop: 59
Location/Qualifiers
                                                                                                                                                                                                                                     Class: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                       Contact: Robert B. University of Utah University of Utah
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly
,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A.
and Wright,D.,Weiss,R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AZ442276 59 bp DNA linear GSS 0 1M0234P19R Mouse 10kb plasmid UUGCIM library Mus musculus clone UUGCIM0234P19 R, DNA sequence.
                                                                                                                                                                                                                                                        Seq primer: CACACAGGAAACAGCTATGACC
                                                                                                                                                                                                                                                                        Plate: 0234
                                                                                                                                                                                                                                                                                                                        Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                        Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AZ442276.1 GI:10566289 GSS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              9 TCCTCATCTGGGGCGGGGGGGGGCCCCCCCAAAAGAGGGGGGGAGA 54
                                                                                                                                                                                                                                                                                                                                                                                                                                                           plasmid inserts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Dunn, D., Aoyagi, A.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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                                                                                                                                                                                                                                                                                  ddunn@genetics.utah.edu
Length: 10000 Std Erro
                 /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"/note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson Laboratory Mouse DNA Resource
                                                                                             /db_xref="taxon:10090"
/clone="UUGC1M0234P19"
/clone_lib="Mouse 10kb plasmid UUGC1M library"
                                                                                                                                                                                                                                                                                                                                                                       Biomedical Polymers Research Bldg.,
                                                                                                                                                     /organism="Mus musculus
/strain="C57BL/6J"
                                                                                     /sex="Male"
http://www.jax.org/resources/documents/dnares/). The DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /db_xrefr"taxon:9606"
/clone="IMAGE:1659302"
/clone_lib="Soares_total_fetus_Nb2HF8_9w"
/dev_stage="8-9_weeks"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /lab_host="DH10B"
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Pred. No. 1.6e+05;
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olumn: 19
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AI722821
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VERSION
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ORIGIN
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        15 AATTTCACCAGTCACCAAGTCCTCAGTACACCTGTGAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Unpublished (1998)
Other_BSTs: fc32all.xl
Contact: Stephen L. Johnson
Washington University School of Medicine
4444 Forest Park Parkway, Box 8501, St. I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Danio rerio
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Actinopterygii; Neopterygii; Teleostei; Ostariophysi; Cypriniformes
                                                                                                                                                                                                                                                         Missouri (web address www.genomesystems.com) (email contact: info@genomesystems.com) and Research Genetics, Huntsville, Alabama (web address: www.resgen.com) (email contact: info@resgen.com) and RessourcenZentrumprimarDatenbank, Berlin, Germany (web address:
                                                                                                                                                                                                                                                                                                                                                Matthew Clark. DNA Sequencing by: Washington University Genome Sequencing Center Clone distribution: Genome Systems, St. Louis, Missouri (web address: www.genomesystems.com) (email contact:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Clark,M., Johnson,S.L., Lehrach,H., Lee,R., Li,F., Marra,M., Eddy,S., Hillier,L., Kucaba,T., Martin,J., Beck,C., Wylie,T., Underwood,K., Steptoe,M., Theising,B., Allen,M., Bowers,Y., Person,B., Swaller,T., Gibbons,M., Pape,D., Harvey,N., Schurk,R., Ritter,E.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AI722821 34 bp mRNA linear EST 07-JUN-2001 fc32all.yl Zebrafish WashU MPIMG EST Danio rerio cDNA clone fmAGE:3723068 5' similar to TR:075840 075840 UBIQUITOUS KRUPPEL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WashU Zebrafish EST Project 1998
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Kohn, S., Shin, T., Jackson, Y., Cardenas, M., McCann, R.,
                                                                                                                                            Trace considered overall poor quality Possible reversed clone: similarity of Seq primer: T3 ET from Amersham
                                                                                                                                                                                                                                                                                                                                                                                                                                                 cDNA Library Preparation: Matthew Clark. cDNA Library Arrayed by:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AI722821.1
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                                                                                                                   High quality sequence stop: 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Email: zbrafish@watson.wustl.edu
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/organism="Danio rerio"
/db_xref="taxon:7955"
                                                                                       location/Qualifiers
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                                                                                                                                                                           wrong strand
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SOURCE KEYWORDS

COMMENT

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RESULT 30
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                                      FEATURES
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            26;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WashU-HHMI Mouse EST Project
Washington University School of MedicineP
4444 Forest Park Parkway, Box 8501, St. LA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 CCTGAAGGCACACCAAAGGACACATACAGGTGA 33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Contact: Marra M/Mouse EST Project
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The WashU-HHMI Mouse EST Project
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Marra, M., Hillier, L., Allen, M., Bowles, M., Dietrich, N., Dub
Geisel, S., Kucaba, T., Lacy, M., Le, M., Martin, J., Morris, M.,
Schellenberg, K., Steptoe, M., Tan, F., Underwood, K., Moore, B.
Theising, B., Wylie, T., Lennon, G., Soares, B., Wilson, R. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus 1 (bases 1 to 58)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mus musculus
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                                                                    High quality sequence stop:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Unpublished (1996)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Waterston, R
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AA139470.1 GI:1701736
                                                                                                               Seq primer: -28m13 rev1 ET from Amersham
                                                                                                                                                          Possible reversed clone: similarity on wrong strand
                                                                                                                                                                                                                                                MGI:369410
                                                                                                                                                                                                                                                                                                                    This clone is available royalty-free through LLNL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        house mouse
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity
                                                                                                                                                                                                                                                                                  IMAGE Consortium (info@image.llnl.gov) for
                                                                                                                                                                                                                                                                                                                                                                                                              314 286 1800
314 286 1810
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    sites of the pSPORTI vector (BRL). Library was constructed by Matthew Clark (Lehrach lab: ICRF, London and Max Planck Institut fuer Molekulare Genetik, Berlin). cDNAs for EST analysis were selected following oligonucleotide hybridization fingerprinting of arrayed clones from zebrafish late somitogenesis (26 ss), adult liver or embryonic shield stage (5.6 h) libraries. Fingerprint data were used to computationally cluster cDNAs, and a single cDNA from each cluster was chosen for sequencing. In some cases multiple members of the same cluster were sequenced to assess clustering parameters or single clones
                                                                                                                                                                                                 considered overall poor quality
                                                                                                                                                                                                                                                                                                                                                                    mouseest@watson.wustl.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             were sequenced additional times to assess quality control."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            double-stranded cDNA was ligated to Sal I adaptors (BRL), digested with Not I and cloned into the Not I and Sal I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   stage embryos"
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                                      Location/Qualifiers
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Pred. No. 1.3e+05;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      56
                                                                                                                                                                                                                 Source: IMAGE Consortium, LLNL
This clone is available royalty-free through LLNL;
IMAGE Consortium (info@image.llnl.gov) for further j
Trace considered overall poor quality
                                                                                                                                                                                                                                                                           High quality sequence starts: 1
High quality sequence stops: 1
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Yi31911.rl Soares placenta
YMACE:140900 5' similar to
                                                                                                                                                                                                                                                                                                                   Email: est@watson.wustl.edu
Insert Size: 1022
                                                                                                                                                                                                                                                                                                                                                   Tel: 314 286 1800
Fax: 314 286 1810
                                                                                                                                                                                                                                                                                                                                                                                 4444 Forest Park Parkway,
                                                                                                                                                                                                                                                                                                                                                                                                                                    Unpublished (1995)
                                                                                                                                                                                                                                                                                                                                                                                                                                                              ,M., Hultman,M., Kucaba,T., Le,M., Lennon,G., Marra,M., Rifkin,L., Rohlfing,T., Soares,M., Tan,F., Trevaskis,E., R., Williamson,A., Wohldmann,P. and Wilson,R.
                                                                                                                                                                                                                                                                                                                                                                                                  Washington University
                                                                                                                                                                                                                                                                                                                                                                                                                      Contact: Wilson RK
                                                                                                                                                                                                                                                                                                                                                                                                                                                   The WashU-Merck EST Project
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Hillier,L., Clark,N., Dubuque,T., Elliston,K.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     30;
                                                                                                                                                                                                   Insert Length:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens
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                                                                                                                                                                                       primer: M13RP1
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                                            /organism="Homo sapiens"
/db_xref="GDB:549524"
/db_xref="taxon:9606"
/clone="IMAGE:140900"
/clone=lib="Soares placenta Nb2HP"
/dev_stage="placenta obtained at birth (full term)"
/lab_host="DH10B (ampicillin resistant)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          'sex="Female"
                                                                                                                                                   Location/Qualifiers
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/strain="NIH Swiss"
/db_xref="Taxon:10090"
/clone="IMAGE:603978"
/clone_lib="Stratagene mouse heart (#937316)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /lab_host="SOLR (kanamycin resistant)"
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/dev_stage="13 day embryos"
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                                                                                                                                                                                                     Std Error: 0.00
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a Nb2HP
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                                                                                                                                                                                                                                                                                                                                                                               Louis,
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                                                                                                                                                                                                                                                                                                             IMAGE Consortium (info@image.llnl.gov) for fit Trace considered overall poor quality Possible reversed clone: similarity on wrong Possible reversed clone: similarity on wrong Possible reversed clone: polyT not found Insert Length: 1709 Std Error: 0.00 Steg primer: -40M13 fwd. from Amersham
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Tel:
                                                                                                                                                                                                                                                                                                                                                                                                                             This clone is available royalty-free through LLNL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              4444 Forest Park Parkway,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Washington University
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Contact: Wilson RK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; I
Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
1 (bases 1 to 43)
Hiller, L., Lennon, G., Becker, M., Bonaldo, M.F., Chian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AA059057

A3 bp mRNA linear EST zf63bl0.sl Soares retina NZb4HR Homo sapiens cDNA clone IMAGE.381595 3. similar to TR:G868160 G868160 CYS2/HIS2
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Email: est@watson.wustl.edu
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quality sequence stop: 1.
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314 286 1810
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              /lab_host="DH10B (ampicillin resistant)"
/note="Organ: eye; Vector: pT7T3D (Pharmacia) with a modified polylinker; Site_1: Not I; Site_2: Eco RI; 1 strand cDNA was primed with a Not I - oligo(dT) prime
/tissue_type="retina"
/dev_stage="55 year o
                                                                                                                                                     /db_xref="taxon:9606"
/clone="IMAGE:381595"
/clone_lib="Soares retina
                                                                                                                                                                                                                  /organism="Homo sapiens"
/db_xref="GDB:1289852"
                                                                                                                                                                                                                                                                           Location/Qualifiers
                                                                                                                                         /sex≠"male"
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modified polylinker; Site_1: Not I; Site_2: Eco RI; 1st
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80.6%; Pred. No. 2.46
tive 0; Mismatches
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RESULT 33
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Eukaryota; Métazoa; Chordata; Čraniata; Vertebrata; Euteleostomi; Mammalla; Eutherla; Primates; Catarrhin; Hominidae; Homo.

1 (bases 1 to 49)

Hillier,L., Clark,N., Dubuque,T., Elliston,K., Hawkins,M., Holman,M., Kucaba,T., Le,M., Lennon,G., Marra,M., Parsons,J., Rifkin,L., Rohlfing,T., Soares,M., Tan,F., Trevaskis,E., Waterston,R., Williamson,A., Wohldman,P. and Wilson,R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              R75644 49 bp mRNA linear EST 06-JUN y159f01.sl Soares placenta Nb2HP Homo sapiens cDNA clone IMAGE: 43545 3' similar to SP:S40468 S40468 PROTEASOME SUBUNIT
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                                                                                                                                                                                                                                                                                                                                                                                    Source: IMAGE Consortium, LLNL
This clone is available royalty-free through LLNL; contact the
IMAGE Consortium (info@image.llnl.gov) for further information.
Trace considered overall poor quality
Possible reversed clone: similarity on wrong strand
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Unpublished (1995)
Contact: Wilson RK
Washington University School of Medicine
4444 Forest Park Parkway, Box 8501, St. 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 High quality sequence starts: 1 High quality sequence stops: 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Tel: 314 286 1800
Fax: 314 286 1810
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RC10-II - ;, mRNA sequence.
R75644
                                                                                                                                                                                                                                                                                                                                                  Insert Size: 848
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Email: est@watson.wustl.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens
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                  /db_xref="taxon:9606"
/clone="IMAGE:143545"
                                                                                                                                                                                                                                       /organism="Homo sapiens"
/db_xref≈"GDB:552718"
                                                                                                                 /dev_stage="placenta obtained at birth (full term)"
/lab host="DH10B (ampicillin resistant)"
                                                                                                                                                                     /clone_lib="Soares placenta Nb2HP"
                                                                                                                                                                                                                                                                                                  Location/Qualifiers
                                                                                                                                                     /sex="Female"
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Pred. No. 2.3e
0; Mismatches
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cDNA was ligated
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to Eco RI adaptors
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RESULT 34
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                                                                                                                                                                                      50 AGAACACAATCAACAGCCGCAGACACTCAGCTATCGATTCCACCCAAAAT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Email: ysuzuki@ims.u-tokyo.ac.jp
Suzuki,Y., Yoshitomo-Nakagawa,K., Maruyama,K., Suyama,A. and Sugano,S. Construction and characterization of a full length-enriched and a 5'-end-enriched cDNA library. Gene 200 (1-2), 149-156 (1997).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Department of Virology
Institute of Medical Science, University of Tokyo
4-6-1, Shirokanedai, Minatoku, Tokyo 108-8639, Japan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Suzuki,Y., Taira,H., Tsunoda,T., Mizushima-Sugano,J.,
H., Ota,T., Isogai,T., Tanaka,T., Morishita,S., Okub
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 50
AU105724 Sugano Homo sapiens
KAT10272, mRNA sequence.
AU105724
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21270072
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 50)
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                                 в02962.1
                                                                                     cSRL-164B7-u cSRL flow sorted
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                                                                                                                                                                                                                                                        Similarity 64.(
32; Conservative
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                                                                  genomic clone cSRL-164B7,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (Pharmacia), digested with Not I and cloned into the Not I and Eco RI sites of the modified pT73 vector. Library went through one round of normalization. Library constructed by Bento Soares and M.Fatima Bonaldo. " 12 c 10 g 14 t 2 others
                                                                                                                                                                                                                                                                                                                                                           /clone_lib="Sugano Homo sapiens cDNA library" /note="Differential display comparison of untreated dimethylfumarate treated U937 cells"
                                                                                                                                                                                                                                                                                                                                                                                                             /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="KAT10272"
                                   GI:1412240
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                                                                                                                                                                                                                                                                          0.8%;
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Pred. No. 2.6e+05;
0; Mismatches 10
                                                                                                                                                                                                                                                        Score 21.2; DB 9;
Pred. No. 2.6e+05;
0; Mismatches 18
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COMMENT
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AW692135
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                     Contact: Plant Biology Division
Plant Biology Division
The Samuel Roberts Noble Foundation
The Parkway, Ardmore, OK
                                                                                                                                                                                         Не, X.-Z.
, C.J., F
                                                                                   On Apr 14, 2000 this sequence version Contact: Dixon RA
                                                                                                                                  Expressed Sequence Tags from the Samuel Medicago truncatula stem library
                                                                                                                                                                                                                                                                                                                                                                                      NF048AllST1F1000 Developing stem Medicago NF048AllST 5', mRNA sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          29;
                                                                                                                       Unpublished (2000)
                                                                                                                                                                                                                                                                           Spermatophyta; Magnoliophyta;
                                                                                                                                                                           , R.A
                                                                                                                                                                                                                                              Medicago
                                                                                                                                                                                                                                                           Rosidae; eurosids I;
                                                                                                                                                                                                                                                                                                           Medicago truncatula
                                                                                                                                                                                                                                                                                                                                 barrel medic.
                                                                                                                                                                                                                                                                                                                                                               AW692135.2 GI:11934083
                                                                                                                                                                                                                                                                                         Eukaryota; Viridiplantae; Streptophyta;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               High quality sequence stop:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 McDermott Center for Human Growth and Development University of Texas Southwestern Medical Center At Dallas 5323 Harry Hines Blvd, Dallas TX 75235-8591
Tel: 214-648-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Evans,G.A., Burbee,D., Davies,C., Hahner,L., Oliver,T., Gilbert,M. Jones,D., Ward,T., Gillilan,E., Schagemann,J., Probst,S.., Harris,J., DeFord,J., McFarland,J., Burzinski,K., Khan,M., Kupfer,K. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Class: cosmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Seq primer: T7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Email: gevans@utsw.swmed.edu, shane@mcdermott.swmed.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Contact: Evans GA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Genomic Sequence Sampled Map of Chromosome 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Garner, H.R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Unpublished
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Sam Noble Parkway,
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580 221 7380
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                                                                                                                                                                                     Flores, H.R.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /note="Vector: sCos-1; Human Chromosome 11 specific cosmid
library prepared from flow sorted human Chromosome 11
derived from Chinese Hampster Ovary (CHO) monochromosomal
somatic cell hybrid, JI"
18 c 6 g 13 t 1 others
                                                                                                                                                                                                       Shadle, G.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="cSRL-164B7"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /cell_type="chimeric hamster somatic cell hybrid"
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                                                                                                                                                                                       Inman,J.T.,
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                                                                                                                                                                                                     Scott, A.D.,
                                                                                                                                                                                                                                                           Fabales;
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Pred. No. 2.6e
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                                                                                                                                                                                                                                                                      core eudicots;
                                                                                                                                              Noble Foundation
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                                                                                                                                                                                                                                                                                    Tracheophyta;
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AZ595836/c
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                                                                                                                                                                                                                                                                         Plate: 0408 row: O column: 08
Seg primer: CACACAGGAAACAGCTATGACC
Class: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                                  Contact: Robert B. University of Utah University of Utah
                                                                                                                                                                                                                                                       High quality sequence stop:
                                                                                                                                                                                                                                                                                                                                                                 Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                     Rm.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A.
                                                                                                                                                                                                                                                                                                                                                   :Trema
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          plasmid inserts
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AZ595836
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Mammalia; Eutheria; Rodentia; Sclurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1M0408008R Mouse 10kb plasmid UUGC1M clone UUGC1M0408008 R, DNA sequence.
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ert Length: 923 Std Error: 0.0
te: 048 row: A column: 11
primer: TCACACAGGAAACAGCTATGAC.
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Length: 10000 std Erro
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                                 /note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson Laboratory Mouse DNA Resource
                (http://www.jax.org/resources/documents/dnares/).
                                                                                                                                               /db_xref="taxon:10090"
/clone="UUGC1M0408008"
                                                                                                                                                                                                                                                                                                                                                                                                                     Biomedical
                                                                                      /lab_host="E. Coli strain XL10-Gold, T1-resistant,
                                                                                                              /sex="Male"
                                                                                                                         /clone_lib="Mouse 10kb plasmid UUGC1M library"
                                                                                                                                                                                  /organism="Mus musculus"
/strain="C57BL/6J"
                                                                                                                                                                                                                                      Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /dev_stage="Pooled developmental"
/note="Vector: Lambda Zap; Contains a mixture of
internodal stem segments"
16 c 2 g 20 t
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/db_xref="taxon:3880"
/clone="NF048AllST"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /clone_lib="Developing stem"
/tissue_type="stem"
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hydrodynamically sheared by repeated passage through
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Pred. No. 2.7e+05;
0; Mismatches 13;
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RESULT 39
AU103658
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ORGANISM
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AU256802/c
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                                                                                        40
                                                                                                                                                          24;
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AU256802 3'-directed mouse cDNA library Mus musculus cDNA clone
BED0008995 3', mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                           Email: kkato@bs.aist-nara.ac.jp,
URL:http://love2.aist-nara.ac.jp/BED/index.html
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Contact: Kikuya Kato
Graduate School of Biological Sciences
Nara Institute of Science and Technology
8916-5 Takayama, Ikoma, Nara 630-0101, Ja
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Generation of expressed sequence tags Unpublished (2002)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AU256802
AU256802.1 GI:20320800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Kato, K. and Matoba, R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mus musculus
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27; Conserv
   AU103658
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  house mouse
                                                                                                                                                                         Similarity
                                                                                                          TGTCTTCCCATTTGGTGTGTTTTCCTCCG 2500
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CTGGAAAGAAACAAGCTGCACAGCCCCAGGAAAACAA 1
                                                                                        TGTCGTTCTATTTGGTGGGCTTTCCCCCG
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                                                                                                                                                        Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWDA2 (g4|473214|gb|h8F129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."

8 c 12 g 18 t
                                                                                                                                                                                                                                         /db_xref="taxon:10090"
/clone="BED0008995"
/clone_lib="3'-directed mouse cDNA
/tissue_type="brain"
/note="Vector: pGEM-T-easy"
10 c 14 g 2 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   purified.
                                                                                                                                                                                                                                                                                                                                                /organism="Mus musculus"
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Pred. No.
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2.6e+05;
                                                                                                                                                     ..6e+05;
5;
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   mRNA
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 30-AUG-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         214 TGGCATTCAGGGATGTGGCTGTGTACTTCACCCAGGA 250
                  Contact: Yutaka Suzuki
Contact: Yutaka Suzuki
Department of Virology
Institute of Medical Science, University of Tokyo
4-6-1, Shirokanedai, Minatoku, Tokyo 108-8639, Japan
4-6-1, Shirokanedai, Minatoku, Tokyo 108-8639, Japan
Email: ysuzuki@ims u-tokyo.ac.jp
Suzuki,Y., Yoshitomo-Nakagawa,K., Maruyama,K., Suyama,A. and Sugano
Suzuki,Y., Yoshitomo-Nakagawa,K., Maruyama,K., Suyama,A. and Sugano
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Suzuki, Y., Yoshitomo-Nakagawa, K., Maruyama, K., Suyama, A. and Sugano, S. Construction and characterization of a full length-enriched and a 5'-end-enriched cDNA library. Gene 200 (1-2), 149-156 (1997).
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AU103658
AU103658.1 GI:13553179
EST.
                                                                                                                                                                                          Diverse transcriptional initiation mapping of mRNA start sites EMBO Rep. 2 (5), 388-393 (2001)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AU105956 50 bp mRNA linear EST 30-AUG-200 AU105956 Sugano Homo sapiens cDNA library Homo sapiens cDNA clone INECO029, mRNA sequence AU105956
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27; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Institute of Medical Science, University of Tokyo 4-6-1, Shirokanedai, Minatoku, Tokyo 108-8639, Japan Email: ysuzukidims.u-tokyo.ac.jp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          EMBO Rep. 2 (5), 388-393 (2001) 21270072
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Suzuki,Y., Taira,H., Tsunoda,T., Mizushima-Sugano,J.,
.H., Ota,T., Isogai,T., Tanaka,T., Morishita,S., Okub
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 50)
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                                                                                                                                                                                                                                                                               Suzuki,Y., Taira,H., Tsu,H., Ota,T., Isogai,T.,
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                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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S. Construction and characterization of a full length-enriched 5'-end-enriched cDNA library. Gene 200 (1-2), 149-156 (1997).
                                                                                                                                                                                                                                                                 Y., Nakamura,Y., Suyama,A. and Sugano,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TGACATTTAGGGATGTGGCCATAGAATTCTCTCTGGA 50
                                                                                                                                                                                                                                                                                                                            (bases 1 to 50)
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/note="Differential display comparison of untreated dimethylfumarate treated U937 cells"
7 c 13 g 14 t 5 others
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="HEP03969"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0.8%;
                                                                                                                                                                                                                                                                                   Tsunoda, T., Mizushima-Sugano, J., T., Tanaka, T., Morishita, S., Okub
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3e+05;
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                                                                                                                                                                                                                                                                                                                                                                            Euteleostomi;
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AA976008/c
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                                                                                                                                                                                                                                                                                                                                Trace considered overall poor quality Insert Length: 453 Std Error: 0.00 Seq primer: -40ml3 fwd. Er from Amersham High quality sequence stop: 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      50
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Christopher A. Moskaluk, M.D.,
Emmert-Buck, M.D., Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AA976008 58 bp mRNA linear oq29b04.s1 NCI_CGAP_:GC4 Homo sapiens cDNA clone IMAC similar to SW:ZN27_HUMAN p17033 ZINC FINGER PROTEIN
                                                                                                                                                                                                                                                                                                                                                                                                                www-bio.llnl.gov/bbrp/image/image.html
                                                                                                                                                                                                                                                                                                                                                                                                                            cDNA Library Arrayed by: Greg Lennon, Ph.D.
DNA Sequencing by: Washington University Genome Sequencing Center Clone distribution: NCI-CGAP clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Unpublished (1997)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    27; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AA976008
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             cDNA Library Preparation: M. Bento Soares, Ph.D.
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                                         /note="Vector: pf7f3D-Pac (Pharmacia) with a modified polylinker; 1st strand cDNA was prepared from 3 pooled germ cell tumors, and was then primed with a Not I oligo(dT) primer. Double-stranded cDNA was ligated to Eco RI adaptors (Pharmacia), digested with Not I and cloned into the Not I and Eco RI sites of the modified pf7f3 vector. Library is normalized. Library was constructed by Bento Soares and M. Fatima Bonaldo."
                                                                                                                                                                                                                              /db_xref="taxon:9606"
/clone="IMAGE:1587727"
/clone_1ib="NCI_CGAP_GC4"
                                                                                                                                                                                                 /tissue_type="pooled germ cell tumors"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                 /organism="Homo sapiens"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /organism="Homo sapiens"
/db_xref="taxon:9606"
/db_xref="taxon:9606"
/clone="InEC0029"
/clone_Tib="Sugano Homo sapiens cDNA library"
/note="Differential display comparison of untreated and dimethylfumarate treated U937 cells"
a 8 c 17 g 12 t
                                                                                                                                                                                                                                                                                                                Location/Qualifiers
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Pred.
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21;
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3e+05;
DΒ
9;
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IMAGE:1587727 3
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                                                                                                                                                                                                                                                                                          RESULT 43
AZ666147/c
LOCUS
                                                                                                                                                                           SOURCE
ORGANISM
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VERSION
                                                                                                                            REFERENCE
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AV957851
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TITLE
                                                                                                           AUTHORS
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Best Local Similarity
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Matches
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                                                                                                                                                                           Mus musculus
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                                                                                                                                                                                                                                            A2666147
                                                                                                                                                                                                                                                                                                                                                                                                                                            28;
                                                                                                                                                                                        house mouse.
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33; Conserv
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Length 58

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COMMENT
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                plasmid inserts
Unpublished (2000)
                                                                                 Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Dunslam,H., Longacre,S., Mahmoud,M., Meenen,E.,
                                                                                                                                     Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus 1 (bases 1 to 56)
                                                Mouse whole genome scaffolding with paired
                                                                      and Wright, D., Weiss, R.
                                                                                                                                                                                                                                                                                  1M0548C04F Mouse 10kb plasmid UUGC1M clone UUGC1M0548C04 F, DNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sakyo-ku, Kyoto, Kyo
Tel: 81-75-753-4081
Fax: 81-75-705-1113
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Satoh, N., Satou, Y., Kohara, Y. and Shin-i, T. Expressed genes in Ciona intestinalis Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Department of Zoology
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ciona intestinalis
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AV957851 Nori Satoh unpublished cDNA library, egg Ciona intestinalis cDNA clone cieg09a19 5', mRNA sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa; Chordata; Urochordata;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Ciona intestinalis.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GGGGAGAAGCCATTTGTATGTGCTGAGTGTGGACGAGGCTTTAATGATAAGTC 1729
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      satoh@ascidian.zool.kyoto-u.ac.jp.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /clone_lib="Nori Satoh unpublished
/tissue_type="whole animal"
/dev_stage="egg"
/note="Vector: pBluescript SK"
/note="Vector: pBluescript SK"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /clone="cieg09a19"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /organism="Ciona intestinalis"
/db_xref="taxon:7719"
                                                                                                                                                                                                                                                    GI:11803293
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); Mismatches 20;
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No. 3.
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3.4e+05;
nes 12;
                                                                                                                                                                                                                                                                                                   library Mus musculus
                                                                                                                    Duval, B., Hamil, C.,
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                                                  end reads from 10kb
                                                                                                                                                                                                                                                                                                                       linear
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                                                                                                    Pedersen, T., Reilly
                                                                                     Niederhausern, A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      49;
                                                                                                                                                                                                                                                                                                                       GSS 14-DEC-2000
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RESULT 44
AZ566268/c
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ORIGIN
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                                                                                                                   JOURNAL
                                                                                                                                          TITLE
                                                                                                                                                             AUTHORS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error:
Plate: 0548 row: C column: 04
Seg primer: CGTTGTAAAACGACGGCCAGT
Class: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         52
                                                                                         1 (bases 1 to 60)

Carlton, J.M. -R. and Dame, J.B.

The Plasmodium vivax and P. berghei gene sequence tag projects Parasitol. Today (Regul. Ed.) 16 (10), 409 (2000)

Contact: Dame JB
                                                                                                                                                                                                                               malaria parasite P. vivax.
Plasmodium vivax
                                                                                                                                                                                                                                                                              GSS
                                                                                                                                                                                                                                                                                                                                                              AZ566268 60 bp DNA linear GSS 07-MAY-219PvD10 Pv MBN #16 (amplified twice) Plasmodium vivax genomic
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Fax: 801 585 7177
Dept. of Pathobiology, College of Veterinary Medicine University of Florida 2015 SW 23rd Avenue, Bldg 1017, Gainesville, FL 32611, USA Tel: 352 392 4700
                                                                                                                                                                                                                                                                                                                         DNA sequence
AZ566268
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             High quality sequence stop: 56.
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                                                                                                                                                                                                        Eukaryota; Alveolata; Apicomplexa; Haemosporida; Plasmodium
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pwD42 (gil4732114)gblAF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and the complementary to the insert adaptors.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      adaptored vector DNA, and transformed into adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-" /note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Biomedical Polymers Research Bldg.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         purified.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (http://www.jax.org/resources/documents/dnares/).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /organism="Mus musculus"
/strain="C57BL/6J"
/db_xref="taxon:10090"
/clone="UUGC1M0548C04"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /sex="Male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /clone_lib="Mouse 10kb plasmid UUGC1M library"
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Pred.
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No. 3.7e+05;
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                                                                                                                                                                                                                                                                                                                              Tissue Procurement: DCTD/DTP
CDNA Library Preparation: Life Technologies, Inc.
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Class: shotgun
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BG575662
602598728F1 NIH_MGC_87 Homo sapiens cDNA clone
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Seq primer: M13(-20) forward
                                                                                                                                                                                                                               found through the I.M.A.G.E. Consortium/LLNL http://image.llnl.gov Plate: LLAM10577 row: g column: 08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Contact: Robert Strausberg, Ph.D. Email: cgapbs-r@mail.nih.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae
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BG575662.1
                                                                                                                                                               High quality sequence stop:
Location/Qualifiers
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//dev_stage="asexual blood forms"

//lab_host="Saimiri boliviensis"

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//note="vector: Lambda ZAP II (Stratag
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a 8 c 7 g 22 t
                                                            /organism="Homo sapiens"
/db_xref="taxon:9606"
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/clone_lib="NIH_MGC_87"
                               /clone="IMAGE:4707487"
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                                                                                                                                                           l Similarity
29; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                     to give a tight size distribution (
4 kb). The v + i method used for the library construction is described in detail in Smith, H. and Venter, J.C. (Making small insert libraries for whole genome shotgun sequencing projects. Genome Sequencing: A Practical Approach, eds. M. Vaudin and B. Barrell, Oxford University Press, 1999).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     32;
                                                                                                                                                                                                                                                                                                                                                                                             Details of T. brucei sequencing at the Sanger Centre are available at http://www.sanger.ac.uk/Projects/T_brucei/.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Submitted (10-DEC-2000) Trypanosoma brucei genome sequencing project, Sanger Centre, The Wellcome Trust Genome Campus, Hinxton, Cambridge CB10 1SA, E-mail: barrell@sanger.ac.uk and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Rockville, MD. Genomic DNA isolated from a cloned population of Trypanosoma brucei (TREU927/4 GUTat 10.1) was mechanically shear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Constructed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Hall,N., Bowman,S., Lennard,N.J., Doggett,J., Atkin,R., Chillingworth,C., Ormond,D., Harris,B., El-Sayed,N., Hou,L., Melville,S.E., Rajandream,M.A. and Barrell,B.G. Direct Submission
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2 CTCTGGACGGTGGTTTGGACAACACAGTCAGGTCCTGGGGACCTGCGCGAG 52
                                                                                                                                                                                                                                                                                                                                                                                                                                      Email: nelsayed@tigr.org
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           nhl@sanger.ac.uk
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Trypanosoma brucei
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llarity 67.4%;
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/strain="TREU927"
/db_xref="taxon:5691"
/clone="328401"
a 16 c 7 g 1
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                                                                                                                                                                                                                                                                                                                                                                              Location/Qualifiers
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62.7%;
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                                         1407 CACTTAAGGACACACACAGGAGAGAAGCCTTATGTATGCACAGAATGTGGG 1457
2 CACATTAAGATCCAACGTGGAGCGAAACCCTTCACCTGCCCTGAATGTGGG 52
                                                                                           l Similarity
32; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Washington University School of Medicine Washington University School of Medicine Park Parkway, Box 8501, St. 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Clark, M., Lehrach, H., Appel, B., Eisen, J., Johnson, S., Marra, M., Eddy, S., Hillier, L., Allen, M., Bowles, L., Dubuqque, T., Geisel, G., Jost, S., Kucaba, T., Lacy, M., Le, N., Lennon, G., Martin, J., Moore, B., Schellenberg, K., Steptoe, M., Tan, F., Theising, B., White, Y., Wylie, T., Materston, R. and Wilson, R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CLONE id field represents a position identifier on the original CDNA Library preparation plate. CDNA Library Preparation: Matthew Clark. CDNA Library Arrayed by: Matthew Clark. DNA Sequencing by: Washington University Genome Sequencing Center Clone distribution: Genome Systems, St. Louis, and Max Planck Institut fuer Molekulare Genetik, Berlin Tel +49 30 84 13 1235
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Prace considered overall poor quality
Possible reversed clone: similarity on wrong strand
Seq primer: T7 ET from Amersham
High quality sequence stop: 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Steve Johnson Lab Internal ID - P3_284 NOTE - For this library, the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            fallh10.rl Zebrafish ICRFzfls Danio rerio cDNA clone 14H24 5, similar to SW:ZG58_XENLA P18730 GASTRULA ZINC FINGER PROTEIN XLCGF58.1; mRNA sequence.
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Actinopterygii; Neopterygii; ?
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314 286 1810
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                                                                                                                                                                                                 digested with Not I and cloned into the Not I and Sal I sites of the pSPORTI vector (BRL). Library was constructed by Matthew Clark (Lehrach Lab; ICRF, London and Max Planck Institut fuer Molekulare Genetik, Berlin) and was not biochemically normalised. 70,000 clones from this library were arrayed on high density filters and subsequently screened by oligonucleotide hybridization fingerprinting to identify unique or minimally redundant clones for more intensive analysis."
                                                                                                                                                                                                                                                                                                                                                                                                                                  /sex⇔"mixed"
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                                                                                                                 0.7%;
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Pred. No. 4.2e+05;
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JOURNAL COMMENT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BH862808 55 bp DNA linear GSS 05-AUG-200 SALK_090646 Arabidopsis thaliana TDNA insertion lines Arabidopsis thaliana genomic clone SALK_090646, DNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          10010 N. Torrey Pines Rotel: 858 453 4100 x1752 Fax: 858 558 6379
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Salk Institute Genomic Analysis Laboratory (SIGnAL)
The Salk Institute for Biological Studies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Alonso,J.M., Leisse,T.J., Barajas,P., Chen,H., Cheuk,R.,
C., Jeske,A., Karnes,M., Kim,C.J., Parker,H., Prednis,L
, 2),mmerman,J. and Ecker,J.R.
A Sequence-Indexed Library of Insertion Mutations in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BH862808.1 GI:22098134
                                                                                                                      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Mammalia; Eutheria; Primates; Catarrhini; Hominidae, 1 (bases 1 to 58)
                                                                                                                                                                                                                                                                                                                             AI955368
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             This is single pass sequence recovered from the left border of TDNA. This sequence lies within 300 bases of the 5' end of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Contact: Joseph R. Ecker
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Arabidopsis Genome Unpublished (2001)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Arabidopsis thaliana
Contact: Robert Strausberg, Ph.D. Email: cgapbs-r@mail.nih.gov
                                      Tumor Gene Index
Unpublished (1997)
                                                                            NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP).
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                                                                                                                                                                                                                                                                    AI955368
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                                                                                                                                                                                   Homo sapiens
                                                                                                                                                                                                        human.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /clone_lib="Ārabidopsis thaliana TDNN insertion lines"
/note="PCR was performed on Arabidopsis thaliana lines
each of which contains one or more TDNA insertion
elements. The resultant fragment for each line was
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      directly sequenced to determine the genomic sequence at the site of insertion. Details of the protocols used can
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /strain="Columbia
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Pred. No. 4.2e
0; Mismatches
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13 g 20 t
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Prednis, L., Shinn,
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Homo.
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E:2507258 3'
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                                                                                                                                                                                                                                          Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Res
                                                                                                                                                                                                                                                                                                                                                                                    Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weiss,R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      50 bp DNA linear GSS 16-FEB-2011M0564L13F Mouse 10kb plasmid UUGC1M library Mus musculus genomic clone UUGC1M0564L13 F, DNA sequence.
                                   High quality sequence stop:
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus 1 (bases 1 to 50)
                                                                                                 Plate: 0564 row: L column: 13 Seq primer: CGTTGTAAAACGACGGCCAGT
                                                                                                                                                                                 Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GSS
                                                                                Class: plasmid ends
                                                                                                                                     Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Erro
                                                                                                                                                                                                                         84112,
                                                                                                                                                                                                                                                                                                                        Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                plasmid inserts
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GI:12883846

GSS 16-FEB-2001

Murinae; Mus.

Mismatches

19;

Indels

0;

Gaps

0

/organism="Mus musculus"

50

Polymers Research Bldg.,

20

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2030 E.,

SLC, TU

Std Error:

0.00

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cDNA Library Preparation: Life Technologies, Inc. cDNA Library Arrayed by: Greg Lennon, Ph.D. DNA Sequencing by: Washington University Genome Sequencing Center Clone distribution: NCI-CGAP clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Trace considered overall poor quality Seq primer: -40UP from Gibco
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Emmert-Buck, M.D., Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Tissue Procurement: Christopher Moskaluk, M.D.,
                                                                                                                                                                                                                                                                                                                                                                                                                                   quality sequence stop:
Location/Qualifiers
                                                                                                                            Average insert size 1.75 kb. Life Technologies catalog 11538-014"
                                                                                                                                                                              /note="Organ: uterus; Vector: pCMV-SPORT6; Site_1: SalI;
Site_2: NotI; Cloned unidirectionally. Primer: Oligo dT
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/tissue_type="well-differentiated endometrial adenocarcinoma, 7 pooled tumors"
/lab_host="DHIOB"
                                                                                                                                                                                                                                                                                                                               /clone="IMAGE: 2507258"
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Search completed: June 4, 2003, 10:24:23 Job time: 3924 secs
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                                                                                                                                                2652 TCAGCTGCAGTGAGGAGAGAATCACCCTGCTGTTGGAGGTTGGGGA 2697
                                                                                                         (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with TA DNA polymerase and TA polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pwD42 (gi14732114|gb haff129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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/note="Vector: PWD42nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackson
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/sex="Male"
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|   | c 225<br>c 226<br>227<br>228   |   | c 219<br>220<br>c 221                           | 217<br>218           |                      | C 213                            |                      | c 208                |                                      |                      | c 202<br>c 203                     |                                   |                                  | 196<br>197           | c 194<br>c 195       | c 192<br>193         | 190<br>191                           | 188<br>c 189                             | c 186<br>c 187                           |                        |                                    |                        | 178<br>c 179                       | 177            |                    | c 173        |                                    | 170      |                      | c 166<br>c 167                               | C 164<br>165         |                 | c 161           |                      | c 156<br>157                         |
|   | 19.8<br>19.8<br>19.8   | 19.8<br>19.8<br>19.8  | 19.8<br>19.8                                    | 19.8                 | 19.8                 | 19.8                             | 19.8                 | 19.8                 | 19.8<br>19.8                         | 19.8<br>19.8         | 19.8<br>19.8                       | 19.8                              | 19.8<br>19.8                     | 19.8<br>19.8         | 19.8<br>19.8         | 20                   | 20                                   | 20<br>20                                 | 20<br>20                                 | 20<br>20               | 20<br>20                           | 20<br>20               | 20<br>20                           | 20             | 20                 | 20           | 20<br>20                           | 20       | 20                   | 20.2   | 20.2                 | 20.2            | 20.2            | 20.                  | 20.2                                 |
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|   |  |   |   |                      |                      |                                  |                      |                      |                                      |                      |                                    |                                   |                                  | •                    | _                    |                      |                                      |  |  |                        |                                    |                        |                                    |                |                    |              |                                    |          |                      |  |                      |                 |                 |                      |                                      |
|   | ABN47162<br>ABN49599<br>ABN49864<br>ABN50011                                     | ABN44277<br>ABN45332<br>ABN46381                            | ABN34919<br>ABN35263<br>ABN43641                | ABN32367<br>ABN33529 | AAV75972             | AAX03235<br>AAC13951             | AA1/6566<br>ABL00876 | AAI74524<br>AAI76039 | AAL37049<br>AAL32905                 | AAL30970<br>AAI76271 | ABK40917<br>ABK33341               | AAD20575<br>AAH26689              | AAV28740<br>AAD21116             | ABK48079<br>ABK48080 | AAA38456<br>AAZ93777 | ABN4 854 9           | ABN42034<br>ABN43740                 | ABN37979<br>ABN39834                     | ABN34721<br>ABN37226                     | ABN33716<br>ABN33813   | ABN33264<br>ABN33445               | AAC65685<br>ABN33155   | AAH91974<br>AAZ21943               | AAL27327       | AD35532            | AAQ81434     | AAQ35070<br>AAZ69441               | AAQ12365 | ABL40933             | ABN48331<br>ABN48650                         | ABN45051<br>ABN47576 | ABN42196        | ABN33459        | AAD37838             | AAL30203<br>AAV19098                 |
|   |  |   |   |                      | •                    |                                  |                      |                      |                                      |                      |                                    | •                                 |                                  |                      |                      |                      |                                      |  |  |                        |                                    |                        |                                    |                |                    |              |                                    |          |                      |  |                      |                 |                 |                      |                                      |
|   | Human spliced tra<br>Human spliced tra<br>Human spliced tra<br>Human spliced tra | Human spliced tra<br>Human spliced tra<br>Human spliced tra | Human spliced<br>Human spliced<br>Human spliced | Human spliced        | •                    | PCR primer used Human secreted   |                      |                      |                                      |                      |                                    | •                                 |                                  |                      |                      |                      |                                      | Human spl<br>Human spl                   |  | Human spl<br>Human spl | Human spl<br>Human spl             | Human BPI<br>Human spl | Human inf<br>Fragment              | Human CLO      | Me                 | HBV targe    | Pro                                | Pro      | Hu                   | Hu:<br>Hu:                                   | Hui                  | Human           | Human           | Human FAIL transm    | Human SNP oligor<br>Synthetic Strept |

| o 0   |                            |   |                            |                            |             |                                  |                            |  |   |                            |                            |                           |                           |                           |                           |   |  |                    |                            |                   |  |  |                   |                         |                            |             |                            |                            |                            |                            |                     |                           |                            |                           |  |                           |                     |                                     |                                     |                           |
|---|----------------------------|---|----------------------------|----------------------------|-------------|----------------------------------|----------------------------|--|---|----------------------------|----------------------------|---------------------------|---------------------------|---------------------------|---------------------------|---|--|--------------------|----------------------------|-------------------|--|--|-------------------|-------------------------|----------------------------|-------------|----------------------------|----------------------------|----------------------------|----------------------------|---------------------|---------------------------|----------------------------|---------------------------|--|---------------------------|---------------------|-------------------------------------|-------------------------------------|---------------------------|
| 298<br>299<br>300   |                            | c 294<br>295                              | c 291<br>292               | 289<br>290                 | 287<br>288  | c 285<br>286                     | 283<br>284                 | 281<br>282   | 280   | 278                        |                            | c 275                     |                           |                           |                           |   | c 267  |                    | 200                        | 261               | c 259<br>260                           |  | 0 256             |                         | c 252                      |             | 249<br>250                 | c 248                      | 240<br>247                 |                            | C 244               |                           | 241                        |                           | c 238                                  |                           | 235                 | C 233                               |                                     | c 230                     |
| 19<br>19  | 19                         |   | 19<br>19                   |                            |             |                                  |                            |  |   |                            |                            |                           |                           |                           |                           |   |  |                    |                            |                   |  | 19                                     |                   |                         |                            |             | . 19                       |                            | 19                         | 19                         | 19                  | 19                        | 19                         | 19                        | 19                                     | 19<br>19                  | 19                  | 19                                  | 19                                  | 19                        |
| 444   | 44                         | 444                                       | 44                         | 44                         | 4.4         | 4                                | . 4                        | .44  | . 4.  |                            |                            |                           |                           |                           |                           |   |  |                    | , o c                      |                   |  |  |                   |                         |                            |             |                            |                            |                            |                            |                     |                           |                            |                           |  |                           |                     |                                     |                                     |                           |
| 0.7   | 0.7                        |   |                            |                            |             |                                  |                            |  |   |                            |                            |                           |                           |                           |                           |   |  |                    | 0.7                        | •                 |  |  | ٠.                | •                       |                            |             |                            | •                          |                            | •                          |                     | •                         |                            |                           |  |                           |                     |                                     |                                     |                           |
| 600   | 600                        | 5 5 5<br>5 6 6<br>5 6 6 6                 | 51                         | 551                        | 50          | 45<br>47                         | 41<br>43                   | 33<br>33   | ω c<br>ω F  | 31<br>00                   | 6 o                        | 000                       | თ თ<br>O O                | 60                        | 0 0                       | 60<br>0                                 | 600  | 0,0,0              | 000                        | 60                | ი ი<br>0                               | 60                                     | 000               | 5 G                     | 5 5<br>6 6                 | 56          | ს ს<br>ნი მ                | 5 3                        | 51                         | 51                         | 51                  | 51                        | 51                         | 51                        | 51                                     | 7 L<br>7 U                | 51                  | 4 <del>4</del> 8                    | 47                                  | 47                        |
| 24 A<br>24 A  |                            |   |                            |                            |             |                                  |                            |  |   |                            |                            |                           |                           |                           |                           |   |  |                    |                            |                   |  |  |                   |                         |                            |             |                            |                            |                            |                            |                     |                           |                            |                           |  |                           |                     |                                     |                                     |                           |
| ABN35000<br>ABN35717<br>ABN35782  | BN3358                     | AH7989<br>AT0474                          | A17331<br>A17860           | AL2911                     | AL2806      | AD3363<br>AZ6695                 | AH7370<br>AT0474           | AQ8948<br>BK4921                                       | AQ4671  | BN5952                     | BN5083                     | BN4838                    | BN4696<br>BN4767          | BN4593                    | BN4555                    | BN4505                                  | BN4115   | BN3975             | BN3415                     | BN3375            | BN3242<br>BN3361                       | BN3235                                 | AT3553            | AV2897                  | AT7888<br>AV2259           | AT9183      | AQ5405<br>A05406           | AH2198                     | AH4011<br>AH4018           | АН3996                     | A1/813<br>AH9041    | AI7339                    | AL3315                     | AL3298                    | AL3265                                 | AA7649                    | AA7648              | AF2951<br>AV1870                    | AZ6851                              | AZ6645                    |
| 270   | 3 3 5                      | 71  | 19                         | 10                         | y 0,        | 37                               | 1 5                        | 28   | 12  | 27                         | ă 4                        | , œ i                     | 7 6                       | 56 6                      | 2 57<br>20 7              | 37 11                                   | 71   | 5 65 6             | 200                        | 0                 | 10,                                    | 22 2                                   | 22                | 33                      | 96                         | 88          | 000                        | . 8                        | 8 6                        | 4                          | 2                   | . 33                      | 92                         | 39                        | 80 #                                   | . 0                       | 88                  | ŭΈ                                  |                                     | 5 55                      |
|   |                            |   |                            |                            |             |                                  |                            |  |   |                            |                            |                           |                           |                           |                           |   |  |                    |                            |                   |  |  |                   |                         |                            |             |                            |                            |                            |                            |                     |                           |                            |                           |  |                           |                     |                                     |                                     |                           |
|   |                            |   |                            |                            |             |                                  |                            |  |   |                            |                            |                           |                           |                           |                           |   | •  |                    |                            |                   |  | :                                      |                   |                         |                            |             |                            |                            |                            |                            |                     |                           |                            |                           |  |                           |                     |                                     |                                     |                           |
| Human<br>Human<br>Human   | Humai                      | PCR pr                                    | Huma<br>Huma               | Huma                       | Human       | ZFP Zi                           | Human<br>PCR p             | uma<br>rar   | HIV a   | Huma                       | Huma                       | Human                     | Huma                      | Huma                      | Huma                      | Human<br>Human                          | Human<br>Human   | Huma               | Human                      | Huma              | Huma                                   | Huma                                   | CMV-              | Huma                    | Mono<br>Sian               | DNA         | Human                      | Huma                       | Huma                       | Huma                       | Huma                | Huma                      | Huma                       | Huma                      | Huma                                   | Human                     | Human               | Humai                               | Huma                                | Huma                      |
| n sg  | SE                         | ime                                       | sil<br>sil                 | SNP                        | SNP         | mar<br>Mar                       | n zinc<br>primer           | n  | am  | 0.00                       | ള                          | ្រ                        | o s                       | 5 5                       | 20                        | s s                                     | s s  | S                  | 0 0                        | n spli            | Human splic                            | ne ımn<br>n spli                       | TPO fu            | n MGMT                  | clonal<br>al per           | encodi      |                            | =                          | 5 5                        | _                          | n clor              | n sile                    | n sile                     | n SNP                     | n SNP                                  | n clon                    | n clor              | n Fak<br>Sapi                       | n map-                              | n me                      |
|   | pliced t                   | contain<br>r, IVHP1                       | ent SNP c                  | oligo                      | oligo       | DNA amp                          | ΒË                         |  | ''' <sub>'''</sub>                                  | ced t                      |                            |                           | ced t                     | iced t                    | ced t                     | iced t                                  |  |                    | iced t                     | ed                | oliced t                               | ımmunogıobı<br>spliced trar            | usion             | r with                  | l anti<br>otide            | ing signa   | -2 typ                     | noglo                      | flank                      | flank                      | e cq4               | ent SN                    | of 1do                     | oligo                     | oligo                                  |                           | e cg20              | ens Iq                              | relate                              | פר                        |
| tran<br>tran<br>tran  | tran                       | ini<br>P1,                                | ⊕<br>0 0                   | nuc                        | o r         | mpl                              | н                          | fici<br>35.3   | rob   | ran                        | ran                        | tran                      | ran                       | ran                       | ran                       | tran                                    | ran  | ran                | tran                       | tran              | tran                                   | ran                                    | pri               | ma.                     | sea                        | gna         | o c<br>V                   | buľ                        | ing                        | ing                        | 392                 | C                         | o d                        | nuc                       | onuc                                   | 045                       | 045                 | G F                                 | e d                                 | e d                       |
|   |                            |   |                            |                            |             |                                  |                            |  |   |                            |                            |                           |                           |                           |                           |   |  |                    |                            |                   |  |  |                   |                         |                            |             |                            |                            |                            |                            |                     |                           |                            |                           |  |                           |                     |                                     |                                     |                           |
|   |                            |   |                            |                            |             |                                  |                            |  |   |                            |                            |                           |                           |                           |                           |   |  |                    |                            |                   |  |  |                   |                         |                            |             |                            |                            |                            |                            |                     |                           |                            |                           |  |                           |                     |                                     |                                     |                           |
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| 371<br>372 ·  | 369                        | 000                                       | 364<br>65                  | 2001                       | 360         | 35<br>5<br>5<br>5<br>5<br>5<br>6 | 55<br>57                   | 55.5   | 353   | 551                        | 50 9                       | 348                       | 346                       | 45                        | 43                        | 341<br>342                              | 140  | 38                 | 3 6 6                      | 34                | 2 6                                    | 310                                    | 29                | 127                     | 2 2 2                      | 24          | 222                        | 21                         | 20                         | 18                         | 17                  | 15                        | 14                         | 12                        | 11                                     | 90                        | 80                  | 06                                  | 05                                  | 203                       |
| 19  | 19<br>19                   | 19  | 19.2<br>19.2               |                            |             |                                  |                            |  |   |                            |                            |                           |                           |                           |                           |   |  |                    |                            |                   |  |  |                   |                         |                            |             |                            |                            |                            |                            |                     |                           |                            |                           |  |                           |                     | 19.4                                | 19.4                                | 19.4                      |
| 0.7<br>0.7  | 0.7                        | 0.7                                       | 0.7                        | 0.7                        | 0.7         | 0.7                              | 0.7                        | 0.7<br>0.7   | 0.7   | 0.7<br>7                   | 0.7                        | 0.7                       | 0.7                       | 0.7                       | 0.7                       | 0.7                                     | 0.7  | 0.7                | 0.7                        | 0.7               | 0.7                                    | 0.7                                    | 0.7               | 0.7                     | 0.7                        |             | 0.7                        |                            |                            |                            | 0.7                 |                           |                            |                           |  | 0.7                       |                     |                                     | 0.7                                 |                           |
|   |                            |   |                            |                            |             |                                  |                            |  |   |                            |                            |                           |                           |                           |                           |   |  |                    |                            |                   |  |  | on a              | <u> </u>                | 44                         | 4           |                            |                            |                            |                            |                     | 2 1                       | <u>თ</u> ტ                 | 6                         | თ თ                                    | , o                       | <u>ه</u> د          | <u>ა</u> თ                          | <u></u>                             | 5 6                       |
| u a a.  | ധധ                         | uωN                                       | തത                         | თ თ o                      | 1010        | מ מ כ                            | 9                          | 9  | 00  | n on o                     | თ თ                        | ارض                       | ש פע                      | on 0                      | יסי                       | ف ف                                     | ق ق  | ເຫຼວ               | ı vi u                     | n (n i            | ıσ                                     | O, C                                   | -                 |                         |                            |             | ~ ~                        | 4                          | 44                         | 4                          | ~ ~                 |                           |                            |                           |  |                           |                     |                                     |                                     |                           |
|   |                            |   |                            |                            |             |                                  |                            |  |   |                            |                            |                           |                           |                           |                           |   |  |                    |                            |                   |  |  |                   |                         |                            |             |                            |                            |                            |                            |                     |                           |                            |                           |  |                           |                     |                                     |                                     | 24                        |
| 17<br>22<br>24  | 24<br>21                   | 22<br>22<br>3                             | 24<br>24                   | 24                         | 2 2 C       | 24                               | 24<br>24                   | 24<br>24   | 24  | 100                        | 24                         | 24                        | 24                        | 24                        | 221                       | 24                                      | 24   | 15                 | 222                        | 22                | 2<br>2<br>2<br>2                       | 22                                     | 22                | 24                      | 220                        | 18          | 15<br>15                   | 22                         | 17<br>22                   | 17                         | 17                  | 24                        | 24                         | 24                        | 24                                     | 24                        | 24                  | 24                                  | 24                                  |                           |
| 17<br>22<br>24  | 24<br>21                   | 22<br>22<br>3                             | 24<br>24                   | 24                         | 2 2 C       | 24                               | 24<br>24                   | 24<br>24   | 24  | 100                        | 24                         | 24                        | 24                        | 24                        | 221                       | 24                                      | 24   | 15                 | 222                        | 22                | 2<br>2<br>2<br>2                       | 22                                     | 22                | 24                      | 220                        | 18          | 15<br>15                   | 22                         | 17<br>22                   | 17                         | 17                  | 24                        | 24                         | 24                        | 24                                     | 24                        | 24                  | 24                                  | 24                                  |                           |
| 17<br>22<br>24  | 24<br>21                   | 22<br>22<br>3                             | 24<br>24                   | 24                         | 2 2 C       | 24                               | 24<br>24                   | 24<br>24   | 24  | 100                        | 24                         | 24                        | 24                        | 24                        | 221                       | 24                                      | 24   | 15                 | 222                        | 22                | 2<br>2<br>2<br>2                       | 22                                     | 22                | 24                      | 220                        | 18          | 15<br>15                   | 22                         | 17<br>22                   | 17                         | 17                  | 24                        | 24                         | 24                        | 24                                     | 24                        | 24                  | 24                                  | 24                                  |                           |
| 17<br>22<br>24  | 24<br>21                   | 22<br>22<br>3                             | 24<br>24                   | 24                         | 2 2 C       | 24                               | 24<br>24                   | 24<br>24   | 24  | 100                        | 24                         | 24                        | 24                        | 24                        | 221                       | 24                                      | 24   | 15                 | 222                        | 22                | 2<br>2<br>2<br>2                       | 22                                     | 22                | 24                      | 220                        | 18          | 15<br>15                   | 22                         | 17<br>22                   | 17                         | 17                  | 24                        | 24                         | 24                        | 24                                     | 24                        | 24                  | 24                                  | 24                                  |                           |
| 17 AAT81668<br>22 AAH46286<br>24 ABK97476   | 24 ABA04802<br>21 AAA12852 | 19 AAZ25858<br>22 AAS13044<br>23 AAC86660 | 24 ABN50134<br>24 ABN50892 | 24 ABN48929<br>24 ABN49309 | 24 ABN48232 | 24 ABN45257<br>24 ABN45601       | 24 ABN42108<br>24 ABN44933 | 24 ABN41701<br>24 ABN42055                             | 24 ABN4U261<br>24 ABN41530                          | 24 ABN39954<br>24 ABN39954 | 24 ABN39187<br>24 ABN39806 | 24 ABN39058               | 24 ABN37602               | 24 ABN37324               | 24 ABN34916               | 24 ABN34159<br>24 ABN34323              | 24 ABN32692<br>24 ABN33855                                 | 15 AAQ58535        | 22 AAI75704<br>22 AAI76788 | 22 AAI74805       | 22 AAL29526<br>22 AAT74804             | 22 AAL34007<br>22 AAL27674             | 22 AAL28266       | 24 ABK82591             | 20 AAX17100<br>22 AAD08040 | 18 AAT63812 | 22 AAH45693<br>15 AAO69350 | 22 AAH45692                | 17 AAT36971<br>22 AAH76683 | 17 AAT36970                | 17 AAT 36954        | 24 ABN14635               | 24 ABN48714<br>24 ABN49482 | 24 ABN47493               | 24 ABN44839<br>24 ABN47038             | 24 ABN41855               | 24 ABN41764         | 24 ABN40449<br>24 ABN41414          | 24 ABN39971<br>24 ABN39908          | ABN39534                  |
| 17 AAT81668<br>22 AAH46286<br>24 ABK97476   | 24 ABA04802<br>21 AAA12852 | 19 AAZ25858<br>22 AAS13044<br>23 AAC86660 | 24 ABN50134<br>24 ABN50892 | 24 ABN48929<br>24 ABN49309 | 24 ABN48232 | 24 ABN45257<br>24 ABN45601       | 24 ABN42108<br>24 ABN44933 | 24 ABN41701 Human<br>24 ABN42055 Human                 | 24 ABN4U261 Human 24 ABN41530 Human                 | 24 ABN39954 Human          | 24 ABN39187 Human          | 24 ABN39058 Human         | 24 ABN37602 Human         | 24 ABN37324 Human         | 24 ABN34916 Human         | 24 ABN34159 Human 24 ABN34323 Human     | 24 ABN32692 Human<br>24 ABN33855 Human                     | 15 AAQ58535 Sequen | 22 AAI76788 Human          | 22 AAI74805 Human | 22 AAL29526 Human 22 AAL74804 Human    | 22 AAL34007 Human<br>22 AAL27674 Human | 22 AAL28266 Human | 24 ABK82591 DNA         | 20 AAX17100<br>22 AAD08040 | 18 AAT63812 | 22 AAH45693<br>15 AAO69350 | 22 AAH45692                | 17 AAT36971<br>22 AAH76683 | 17 AAT36970                | 17 AAT 36954        | 24 ABN14635               | 24 ABN48714<br>24 ABN49482 | 24 ABN47493 Human         | 24 ABN44839 Human<br>24 ABN47038 Human | 24 ABN41855 Human         | 24 ABN41764 Human   | 24 ABN40449 Human 24 ABN41414 Himan | 24 ABN39371 Human 24 ABN39908 Human | ABN39534 Human            |
| 17 AAT81668<br>22 AAH46286<br>24 ABK97476   | 24 ABA04802<br>21 AAA12852 | 19 AAZ25858<br>22 AAS13044<br>23 AAC86660 | 24 ABN50134<br>24 ABN50892 | 24 ABN48929<br>24 ABN49309 | 24 ABN48232 | 24 ABN45257<br>24 ABN45601       | 24 ABN42108<br>24 ABN44933 | 24 ABN41701 Human: 24 ABN42055 Human:                  | 24 ABN41530 Human :                                 | 24 ABN39954 Human          | 24 ABN39187 Human :        | 24 ABN39058 Human :       | 24 ABN37602 Human :       | 24 ABN37324 Human :       | 24 ABN34916 Human         | 24 ABN34159 Human : 24 ABN34323 Human : | 24 ABN32692 Human : 24 ABN33855 Human :                    | 15 AAQ58535 Sequen | 22 AAI76788 Human          | 22 AAI74805 Human | 22 AAL29526 Human<br>22 AAL74804 Human | 22 AAL34007 Human<br>22 AAL27674 Human | 22 AAL28266 Human | 24 ABK82591 DNA         | 20 AAX17100<br>22 AAD08040 | 18 AAT63812 | 22 AAH45693<br>15 AAO69350 | 22 AAH45692                | 17 AAT36971<br>22 AAH76683 | 17 AAT36970                | 17 AAT 36954        | 24 ABN14635               | 24 ABN48714<br>24 ABN49482 | 24 ABN47493 Human s       | 24 ABN44839 Human s                    | 24 ABN41855 Human s       | 24 ABN41764 Human s | 24 ABN40449 Human :                 | 24 ABN399/1 Human :                 | ABN39534 Human spl        |
| 38 17 AAT81668 Human c-myb hammer 41 22 AAH46286 Human proliferatin 41 24 ABK97476 Human oestrogen ge | 24 ABA04802<br>21 AAA12852 | 19 AAZ25858<br>22 AAS13044<br>23 AAC86660 | 24 ABN50134<br>24 ABN50892 | 24 ABN48929<br>24 ABN49309 | 24 ABN48232 | 24 ABN45257<br>24 ABN45601       | 24 ABN42108<br>24 ABN44933 | 24 ABN41701 Human spliced<br>24 ABN42055 Human spliced | 24 ABN4U261 Human spliced 24 ABN41530 Human spliced | 24 ABN39954 Human spliced  | 24 ABN39187 Human spliced  | 24 ABN39058 Human spliced | 24 ABN37602 Human spliced | 24 ABN37324 Human spliced | 24 ABN34916 Human spliced | 24 ABN34159 Human spliced t             | 24 ABN32692 Human spliced t<br>24 ABN33855 Human spliced t | 15 AAQ58535 Sequen | 22 AAI76788 Human          | 22 AAI74805 Human | 22 AAL29526 Human<br>22 AAL74804 Human | 22 AAL34007 Human<br>22 AAL27674 Human | 22 AAL28266 Human | 24 ABK82591 DNA binding | 20 AAX17100<br>22 AAD08040 | 18 AAT63812 | 22 AAH45693<br>15 AAO69350 | 22 AAH45692 Human zinc fir | 17 AAT36971 Oligonucleotic | 17 AAT36970 Oligonucleotic | 17 AAT36954 Human C | 24 ABN14635 Human GDMLP-1 | 24 ABN48714 Human spliced  | 24 ABN47493 Human spliced | 24 ABN44839 Human spliced              | 24 ABN41855 Human spliced | 24 ABN41764 Human s | 24 ABN40449 Human spliced           | 24 ABN399/1 Human :                 | ABN39534 Human spliced tr |

| 9 0.7 60 24 ABN5552  | 0.7 60 24<br>0.7 60 24   | 0.7 60 24<br>0.7 60 24<br>0.7 60 24<br>0.7 60 24<br>0.7 60 24<br>0.7 60 24                | 0.7 59 21<br>0.7 60 22<br>0.7 60 22<br>0.7 60 24<br>0.7 60 24<br>0.7 60 24   | 0.7 51 22 0.7 51 22 0.7 51 22 0.7 51 22 0.7 51 22 0.7 51 22 0.7 51 23 0.7 54 17 55 1 | 0.7 51 22 0.7 51 22 0.7 51 22 0.7 51 22 0.7 5 | 0.7 48 24 0.7 48 24 0.7 50 11 50 21 0.7 50 21 0.7 50 21 50 22 0.7   |
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| ALIGNMENTS  ALIGNMENT  ALIGNMENTS  ALIGNMENTS  ALIGNMENTS  ALIGNMENT  ALIGNMENT  ALIGNMENT  ALIGNMENT  ALIGNMENT  ALIGNMENTS  ALIGNMENTS  ALIGNMENTS   | 7 60 244<br>60 244<br>7 60 244<br>7 60 244<br>7 60 244<br>60 244<br>60 244   | . 7 60 24<br>. 7 60 24   | 57 17<br>59 21<br>.7 60 22<br>.7 60 24<br>.7 60 24<br>.7 60 24<br>.7 60 24   | .7 51 22 .7 51 22 .7 51 22 .7 51 22 .7 51 22 .7 51 22 .7 51 22 .7 54 23 .7 55 27 17 57 57 17   | .7 50 22<br>.7 50 22<br>.7 51 22<br>.7 51 22<br>.7 51 22<br>.7 51 22<br>.7 51 22<br>.7 51 22   | .7 46 22<br>.7 48 24<br>.7 50 11<br>.7 50 20<br>.7 50 21<br>.7 50 21<br>.7 50 21  |
| 24 ABN58522 24 AAD29928 25 AAF75943 26 AAF75943 27 AAF75943 28 AAF75943 29 AAF75943 20 AAF75943 21 AAF75943 22 AAF75943 23 AAF75943 24 ABL40716 25 AAF75943 26 ABSS342 27 AABSS342 28 ABSS342 29 AAF85932 20 AAF86003 21 AAZ65003 21 AAZ66003 21 AAZ66003 21 AAZ66003 21 AAZ66003 21 AAZ66003 21 AAZ66003 21 AAZ66084 21 AAZ66003 21 AAZ66084 21 AAZ66003 22 AAL39798 32 Human map-related 33 AAQ35798 34 Human map-related 34 AAC35798 35 Human map-related 36 AAC35798 36 Human map-related 37 AAC35798 38 Human map-related 39 AAC35798 39 Human map-related 30 AAC35798 30 Human map-related 31 AAZ68911 32 AAL39708 33 Human map-related 34 AAC35798 34 Human map-related 35 Human map-related 36 AAC35798 36 Human map-related 37 AAC35798 38 Human map-related 38 Human map-related 39 AAC35798 40 Human map-related 40 AX 40 Human spliced transcript detection oligonucleotide lotion; 40 ABC8911 41 ABN34034 42 Standard; DNA; 60 BP. 42 ABN34034 43 Standard; DNA; 60 BP. 44 ABN34034 4 Standard; DNA; 60 BP. 46 BP. 46 BP. 47 ABN34034 4 Standard; DNA; 60 BP. 48 ABN34034 4 Standard; DNA; 60 BP. 4 ABN34034 4 Standard; DNA; 60 BP.  | 1 C C C C C C C C C C C C C C C C C C C  | 2000000000  | 00000000000000000000000000000000000000   | 22<br>22<br>22<br>23<br>21<br>21   | 2222222222   | 24<br>24<br>20<br>21<br>21<br>21  |
| ABNS8522  ABNS9528  ANTOP928  ANTOP928  ANTOP928  ANTOP928  ANTOP928  ANTOP929  ANTOPP29  ANTOPP |  |   |  |  |  |   |
| Human spliced tran Oligonucleotide us Human single nucle TMV-based randon 1 Oligonucleotide #1 Human myosin heavy  XX Human map-related Human map-related Human map-related Human map-related Human map-related Human map-related Human mp-related Human mp-related Human mp-related  XX Human spliced transcript detection oligonucleoti XX Human SNP oligonuc   | ABN44449 ABN45251 ABN46757 ABN46802 ABN46969 ABN46969 ABN46995 ABN46995 ABN47943 ABN47943 ABN47943 ABN47963  | ABN 3684 ABN 3684 ABN 3684 ABN 38197 ABN 39382 ABN 40173 ABN 40173 ABN 42944 ABN 42944    | AAY128358<br>AAA296824<br>AAF31238<br>ABN33712<br>ABN33743<br>ABN34130<br>ABN34130<br>ABN34393<br>ABN35343   | AAI74758<br>AAI76764<br>AAI78672<br>AAI79706<br>AAH79607<br>AAH39616<br>ABL00455<br>AAC29085<br>AAC29085   | AAF72<br>AAF72<br>AAC97<br>AAL27<br>AAL28<br>AAL28<br>AAL28<br>AAL28<br>AAL29<br>AAL33   |   |
| RESULT 1 ABN34034 ID ABN34034 standard; DNA; 60 BP. XX AC ABN34034; XX DT 15-JUL-2002 (first entry) XX DE Human spliced transcript detection oligonucleotix XX CW Human; mouse; rat; splice transcript; detection; KW splice variant; transcriptome; oligonucleotide loss of the splice transcript of th |  |   |  |  | 10035<br>10035<br>10035  | AAD19782  |
| RESULT 1 ABN34034 ID ABN34034 standard; DNA; 60 BP. XX AC ABN34034; XX AC ABN34034; XX DT 15-JUL-2002 (first entry) XX DE Human spliced transcript detection oligonucleotion; XX KW Human; mouse; rat; splice transcript; detection; KW splice variant; transcriptome; oligonucleotide l XX OS Homo sapiens.   | ** * * * * * * * * * * * *   | <b>=====</b> ===============================  | 5  |  |  | ж О ж Ф Ф Ф ж ::  |
| RESULT 1 ABN34034 ID ABN34034 standard; DNA; 60 BP. XX AC ABN34034; XX AC ABN34034; XX AC ABN34034; XX DT 15-JUL-2002 (first entry) XX DE Human spliced transcript detection oligonucleotion; XX KW Human; mouse; rat; splice transcript; detection; XX XX CS Homo sapiens.  | Human spli<br>Human spli<br>Human spli<br>Human spli<br>Human spli<br>Human spli<br>Human spli<br>Human spli<br>Human spli   | spl   | HCV primer us S. cerevisiae Novel BAC vec Human spliced Human spliced Human spliced Human spliced Human spliced  | Human sile Human sile Human sile Human cons Human DNA Human SNP Human SNP Human sile Primer NOT  | Human PRO pol<br>Human PRO pol<br>Human SNP oll<br>Human SNP oll<br>Human SNP oll<br>Human SNP oll<br>Human SNP oll<br>Human SNP oll   | ST<br>08 p<br>ERG<br>LLA2<br>LLA2<br>Use<br>for<br>for  |
| RESULT 1 ABN34034 ID ABN34034 standard; DNA; 60 BP. XX AC ABN34034; XX AC ABN34034; XX AC ABN34034; XX DT 15-JUL-2002 (first entry) XX DE Human spliced transcript detection oligonucleotion; XX KW Human; mouse; rat; splice transcript; detection; XX XX CS Homo sapiens.  | liced tra-<br>liced tra-<br>liced tra-<br>liced tra-<br>liced tra-<br>liced tra-<br>liced tra-<br>liced tra-<br>liced tra-   | liced tra | mer used in visiae gene AC vector c pliced tran pliced | silent SNP c<br>silent SNP c<br>silent SNP c<br>silent SNP c<br>conservati/e<br>DNA containi<br>SNP flanking<br>silent nonco<br>r NOTKINCYS f  | polypept<br>228 hybri<br>oligonuc<br>oligonuc<br>oligonuc<br>oligonuc<br>oligonuc<br>oligonuc  | r drug-meta primer use RG amberzym RG amberzym A2C.75 ampl A2C.75 ampl Bed to isol FR0228 cD FR0228 cD FR0228 cD  |
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| ALIGNMENTS 4034 standard; DNA; 60 BP. 4034; UL-2002 (first entry) n spliced transcript detection oligonucleotic racing splice transcript; detection; mouse; rat; splice transcript; detection; ce variant; transcriptome; oligonucleotide lespises.  |  |   |  | 0 0 466<br>0 468<br>0 0 471<br>0 0 471<br>1 471<br>1 471<br>1 471  |  | 0 0 0 4 4 4 5 5 6 5 6 5 6 5 6 6 6 6 6 6 6 6 6   |
| ALIGNMENTS  rd; DNA; 60 BP.  irst entry)  irst entry)  transcript detection oligonucleotiat; splice transcript; detection; transcriptome; oligonucleotide l  |  | 18.8<br>18.8<br>18.8<br>18.8<br>18.6<br>18.6<br>18.6                                      | 1888888  | 188  |  |   |
| ALIGNMENTS  NNA; 60 BP.  entry)  rript detection oligonucleotide SEQ  splice transcript; detection; RNA transcriptome; oligonucleotide library;  |  |   |  |  |  |   |
| ALIGNMENTS  60 BP.  detection oligonucleotide SEQ detection; detection; RNA tr. ptome; oligonucleotide library;  |  |   | 00000000   |  |  |   |
| ALIGNMENTS  ction oligonucleotide SEQ inscript; detection; RNA tr coligonucleotide library;  |  |   | 000000000000000000000000000000000000000  |  | . 8 0.7<br>. 8 0.7<br>. 8 0.7<br>. 8 0.7   |   |
| otide SEO<br>On; RNA tr  | .7 36 21<br>.7 39 19<br>.7 42 18<br>.7 45 12<br>.7 45 24<br>.7 46 20<br>.7 47 15   | .7 60 24<br>.7 60 24<br>.7 60 24<br>.7 60 24<br>.7 60 24<br>.7 60 24<br>.7 25 21          | 0.7 60 24<br>0.7 60 24<br>0.7 60 24<br>0.7 60 24<br>0.7 60 24<br>0.7 60 24   | .8 0.7 60 24 .8 0.7 60 24 .8 0.7 60 24 .8 0.7 60 24 .8 0.7 60 24 .8 0.7 60 24 .8 0.7 60 24 .8 0.7 60 24 .8 0.7 60 24 .8 0.7 60 24  | . 8 0.7 53 19<br>. 8 0.7 53 19<br>. 8 0.7 54 22<br>. 8 0.7 56 22<br>. 8 0.7 57 21<br>. 8 0.7 58 21<br>. 8 0.7 59 24<br>. 8 0.7 59 24<br>. 8 0.7 59 24<br>. 8 0.7 60 21   | .8 0.7 50<br>244<br>8 0.7 51<br>22<br>.8 0.7 51<br>.8 0 |
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| NO:6782.<br>script;  | 7 36 16 ANY55895<br>7 38 21 ANAC47336<br>7 39 19 ANV09416<br>7 42 18 ANV04442<br>.7 43 21 ANAC4516<br>7 45 12 ANG12264<br>.7 45 24 ABA04690<br>.7 45 24 ABA04690<br>.7 46 20 ANX19349<br>.7 47 15 ANG68630 | .7 60 24<br>.7 60 24<br>.7 60 24<br>.7 60 24<br>.7 60 24<br>.7 60 24<br>.7 25 21          | 0.7 60 24 ABN37407<br>0.7 60 24 ABN37451<br>0.7 60 24 ABN38146<br>0.7 60 24 ABN38623<br>0.7 60 24 ABN3895<br>0.7 60 24 ABN38931<br>0.7 60 24 ABN40848<br>0.7 60 24 ABN40848  | .8 0.7 60 24 ABN32924  .8 0.7 60 24 ABN33211  .8 0.7 60 24 ABN33872  .8 0.7 60 24 ABN34550  .8 0.7 60 24 ABN34550  .8 0.7 60 24 ABN35020  .8 0.7 60 24 ABN35624  .8 0.7 60 24 ABN35870  | . 8 0.7 53 19<br>. 8 0.7 53 19<br>. 8 0.7 54 22<br>. 8 0.7 56 22<br>. 8 0.7 57 21<br>. 8 0.7 58 21<br>. 8 0.7 59 24<br>. 8 0.7 59 24<br>. 8 0.7 59 24<br>. 8 0.7 60 21   | .8 0.7 50 24 ABL59727  .8 0.7 51 22 AAL29922  .8 0.7 51 22 AAD17720  .8 0.7 51 22 AAJ174658  .8 0.7 51 22 AAJ174658  .8 0.7 51 22 AAJ75286  .8 0.7 51 22 AAJ76883  .8 0.7 51 22 AAJ76883  |

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RESULT 2
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AC ABL004
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DT 05-MAR
DT 05-MAR
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KW Human;
KW inmuno
KW autoim
KW infect
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Best Local
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                                                                                                                                                                                                                                                                                                                                                       quantitatively characterising the corresponding transcriptome, and in detecting RNA transcripts and splice variants of human or animal transcriptomes. The libraries may also be used as specialised mini libraries to detect transcripts of a sub-transcriptome under a particular biological or pathological state, and so allowing the detection of tissue- and pathology-specific genes such as those genes only expressed in specific tissue under a specific pathological condition; to detect developmental specific genes; and to detect RNA transcripts and splice variants of a transcriptome of a patient suffering from a particular disorder. ABN27253 to ABN59589 represent oligonucleotide sequences from rats, humans and mice, which are used in the exemplification of the present invention.
                      Human; single nucleotide polymorphism; SNP; polymorphism; cytostatic; immunosuppressive; antiinflammatory; neuroprotective; antimicrobial; autoimmune disease; inflammation; cancer; nervous system disease;
                                                                                                                                                                                                                                                                                                                      N.B. The sequence data for this patent did not form properties of specification, but was obtained in electronic format at ftp.wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               set of messenger RNAs transcribed from a given transcription unit of the genome, which encodes one or more messenger RNA splice variants. The oligonuclectide libraries are useful for detecting mRNAs from a biological sample, in expression profiling studies, in qualitatively or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New oligonucleotide libraries comprising oligonucleotides which selectively hybridize to mRNAs transcribed from a transcription a genome, useful for detecting tissue-, pathology-, and developmental-specific genes
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                                                                                                                                            ABL00474 standard; DNA;
                                                                                                                                                                                                                                                                                                 Sequence 60
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02-MAY-2001;
                                                                                             05-MAR-2002
                                                                                                                     ABL00474
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present invention describes oligonucleotide libraries for detecting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 1; SEQ ID 6782; 47pp; English
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             infection; polymorphic
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                                                                                                                                                                                                                                                                                                 BP; 16
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2001US-287724P
                                                                      noncoding
                                                                                            (first entry)
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            inflammation; c
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                                                                    SNP oligonucleotide SEQ
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                                                                                                                                                                                                                                                  Mismatches
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hes 0;
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22-NOV-2000; 2000WO-US32311.
                       31-MAY-2001
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RESULT 3
ABL00878/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           antimicrobial activities. Nucleic acids, polypeptides, oligonucleotides and antibodies from the present invention can be used for treating a subject suffering from, at risk for, or suspected of suffering from a pathology ascribed to the presence of a sequence polymorphism. The pathology may be autoimmune diseases, inflammation, cancer, diseases of the nervous system, and infection by pathogenic microorganisms. The SNPs are also useful for determining which forms of a characterised polymorphism are present in individuals. The antibodies may be used in the detection, quantitation and/or cellular or tissue localisation of a polymorphic protein (e.g., for use in measuring levels of the polymorphic protein within appropriate physiological samples).
                                                                                                                                                     Human; single nucleotide polymorphism; SNP; polymorphism; cytostatic; immunosuppressive; antiinflammatory; neuroprotective; antimicrobial;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     oligonucleotides. The sequences from the present invention can
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   comprising one or more single nucleotide polymorphisms (SNPs). to ABB56903 represent human peptides encoded by some of the SN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Isolated human nucleic acids comprising one or more single nucleotide polymorphisms, useful for treating a subject suffering from a pathology, e.g. autoimmune diseases, ascribed to the presence of a sequence polymorphism
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Shimkets RA,
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WO200138586-A2
                                                                                                   infection; polymorphic
                                                                                                                                                                                                                                   Human amino acid change SNP oligonucleotide SEQ
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                                                                                                                                                                                                                                                                                                                                               ABL00878,
                                                                                                                                                                                                                                                                                                                                                                                                    ABL00878 standard; DNA; 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 51 BP; 9 A; 10 C; 14 G; 18 T; 0 other;
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                                                                                                                               disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
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98.0%;
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Pred. No. 2.8e-05;
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                                                                                                                               cancer; nervous
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 23;
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RESULT 4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              comprising one or more single nucleotide polymorphisms (SNPS). ABB56531
Cto ABB56903 represent human peptides encoded by some of the SNP
Coligonucleotides. The sequences from the present invention can have
immunosuppressive, cytostatic, antiinflammatory, neuroprotective and
continicrobial activities. Nucleic acids, polypeptides, oligonucleotides
cand antibodies from the present invention can be used for treating a
continicrobial activities. State for, or suspected of, suffering from a
continicrobial activities of the presence of a sequence polymorphism. The
pathology ascribed to the presence of a sequence polymorphism. The
continion protein for determining which forms of a characterised
continuous system, and infection by pathogenic microorganisms. The sNPs
care also useful for determining which forms of a characterised
continuous present in individuals. The antibodies may be used in
continuous protein (e.g., for use in measuring levels of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
WPI; 2002-257383/30
                                                                                 28-JUL-2000;
02-MAY-2001;
                                                                                                                                                                                                                                                                                 Human spliced transcript detection oligonucleotide SEQ ID NO:6663.
                                                                                                                                                                                                                                                                                                                15-JUL-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                         ABN33915 standard;
                          Shoshan A,
                                                                                                                            20-JUL-2001;
                                                                                                                                                                                     WO200210449-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   polymorphic protein (e.g., for use in measuring levels of the polymorphic protein within appropriate physiological samples).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABL00010 to ABL01104 represent human nucleic acid oligonucleotides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Shimkets
                                                       (COMP-) COMPUGEN INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Isolated human nucleic acids comprising one or more single nucleotide polymorphisms, useful for treating a subject suffering from a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         pathology, e.g. autoimmune diseases, ascribed to the presence of a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         24-NOV-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                          variant;
                                                                                                                                                                                                                                                        mouse; rat; splice transcript; detection; RNA transcript
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity 50; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           51 BP; 10 A; 14 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Page 509; 674pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               polymorphism -
                         Wasserman
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                 2000US-221607P
2001US-287724P
                                                                                                                            2001WO-IB01903
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Leach M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      99US-0167383
                                                                                                                                                                                                                                            transcriptome; oligonucleotide library;
                                                                                                                                                                                                                                                                                                                                                                         DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1.8%;
                          ٦
                                                                                                                                                                                                                                                                                                                                                                         60
                          Mintz
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           12 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 49.4; DB 23
Pred. No. 2.8e-05;
D; Mismatches 1
                          (F)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           15 T; 0 other;
                          Mintz
                         Ļ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              23;
                       Faigler
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
                         s
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               51;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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RESULT 5
AAI77643
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  XXXX
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CC transcription units that populate a genome. The library comprises conserveral oligonucleotides, each capable of hybridising selectively to a certain of messenger RNAs transcribed from a given transcription unit of the genome, which encodes one or more messenger RNA splice variants. CC The oligonucleotide libraries are useful for detecting mRNAs from a certain contracteristing the corresponding transcriptome, and in contracteristing the corresponding transcriptome, and in contracteristing the corresponding transcriptome, and in certain contracteristing the corresponding transcriptome, and in certain contracteristic problems and splice variants of human or animal contracteristic contracterist
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Best Loc
Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                                                                                 Human; single nucleotide polymorphism; SNP; genome; protein therapy; vaccine; probe; diagnostic assay; quantitation; restorative therapy; polymorphic; ds.
                                                                                                       30-NOV-1999;
29-NOV-2000;
                                                                                                                                                                       30-NOV-2000; 2000WO-US32758
                                                                                                                                                                                                                     07-JUN-2001.
                                                                                                                                                                                                                                                              WO200140521-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                            Human silent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      09-NOV-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAI77643;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      messenger RNAs that populate a (sub-)transcriptome, where the (sub-)transcriptome comprises messenger RNAs transcribed from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New oligonucleotide libraries comprising oligonucleotides which selectively hybridize to mRNAs transcribed from a transcription unit a genome, useful for detecting tissue-, pathology-, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAI77643 standard; DNA; 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         developmental-specific genes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQ
                                                                                                       99US-0168138.
2000US-0726173.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BP; 20 A; 19 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    invention describes oligonucleotide libraries
                                                                                                                                                                                                                                                                                                                                                                                                                                               SNP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ID 6663; 47pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                        containing nucleic acid SEQ:4584
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1.3%; Score 35; DB: 74.6%; Pred. No. 1.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              12 G; 9 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 24; Length 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       15;
                                                                                                                                                                                                                                                                                                                                                                          detection
                                                                                                                                                                                                                                                                                                                                                                                          gene therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      multiple
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  detecting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1099
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  59
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Shimkets RA,

Leach M;

(CURA-) CURAGEN CORP

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RESULT 6
AAL26919
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Дb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                polypeptides by expressing inactive proteins or to supplement the patients own production of polypeptide. Additionally, (I) and its complementary sequences may also be used as DNA probes in diagnostic assays to detect and quantitate the presence of similar nucleic acids in samples, and therefore which patients may be in need of restorative therapy. The polypeptides encoded by (I) may be used as antigens in the production of antibodies specific for polymorphic polypeptides. The antibodies may also be used to down regulate expression and activity. The antibodies may also be used as diagnostic agents for detecting the procession of a company of polymorphic and activity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAM53114 to AAM53329 represent peptides related to human polymorphic polynucleotide sequences. The sequences can be used in gene and protein therapy, and in vaccine production. (I) and the polypeptides encoded by them may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate expression of polymorphic polypeptides. For example, (I) may be used to treat discorders by rectifying mutations or deletions in a patient's genome that affect the activity of
                                                                                                                                                                                                                                                                                                                               amyloid protein; angiopoietin; apoptosis related protein; cadherin; cyclin; polymerase; oncogene; histone; kinase; colony stimulating factor; complement related protein; cytochrome; kinesin; cytokine; interferon;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   sequences (I), which AAM53114 to AAM53329
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2001-356160/37
                                                                                              28-DEC-1999;
27-DEC-2000;
                                                                                                                                                                                                                                                   Homo
                                                                                                                                                                                                                                                                                                                                                                                   neuroprotective;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAL26919 standard; DNA; 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 1;
WPI; 2001-465210/50
                                Shimkets
                                                                                                                                                                                   05-JUL-2001
                                                                                                                                                                                                                  WO200147944-A2
                                                                                                                                                                                                                                                                                interleukin; G-protein coupled receptor; thioesterase;
multifactorial disease; autoimmune disease; infection;
nervous system disease; ss.
                                                                                                                                                                                                                                                                                                                                                                               Immunosuppressive; immunostimulatory; antiinflammatory; cytostatic;
neuroprotective; antimicrobial; gene therapy; vaccine; amylase; car
                                                                                                                                                                                                                                                                                                                                                                                                                                   Human SNP oligonucleotide #127
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       presence of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAI73060 to AAI79867
                                                                                                                                                28-DEC-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Polymorphic nucleic acid sequences, useful in genetic testing and
                                                               (CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2375
                                                                                                                                                                                                                                                   sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 Similarity
40; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ш
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AGTCAAATGCCCAGGCAGATAGGGGTGGGTACCTGGTGAAACCCAACCT
                                RA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Page 1913; 2653pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AGACAAATGCCTAGGCAGATAGGGGTAGGTCAACAGTGAAACCCCCACCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      B₽;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     polymorphic polypeptides in samples
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                2000WO-US35498.
                                                                                              2000US-0173419.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                Leach M;
                                                                                                                99US-0173419
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Α,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     represent isolated human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        contain single nucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      14 C; 13 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 34.6; DB Pred. No. 1:4; 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    7 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     9;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     polymorphic polynucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  polymorphisms
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length
                                                                                                                                                                                                                                                                                                                    inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     51;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2423
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (SNPs)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                   cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0;
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RESULT 7
AAI77642
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     G-protein coupled receptors and thioesterases. The present sequence is one such oligonucleotide. The oligonucleotides and the peptides encoded by them may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate expression of the proteins listed above. Disorders that may be prevented, diagnosed and/or treated include multifactorial diseases with a genetic component, such as autoimmune diseases (e.g. rheumatoid arthritis, multiple sclerosis, diabetes, systemic lupus erythromatosus and Grave's disease), inflammation, cancer (e.g. cancers of the bladder, brain, breast, colon and kidney, leukaemia), diseases of the nervous system and an infection of pathogenic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  variants of proteins related to amylases, amyloid proteins, angiopoi apoptosis related proteins, cadherin, cyclin, polymerase, oncogenes, histones, kinases, colony stimulating factors, complement related
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Polymorphic nucleic acids encoding e.g. amylases, oncogenes and histones, useful for diagnosing and cancer, autoimmune diseases and infections -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 1; Page 1434; 4143pp; English.
                                     Claim 1; Page 1913; 2653pp; English
                                                                                                                                                                                                                                                       07-JUN-2001
                                                                                                                                                                                                                                                                                                                                      quantitation;
                                                                                                                                                                                                                                                                                                                                                 protein therapy;
                                                                                                                                                                                                                                                                                                                                                                                        Human silent SNP containing nucleic acid SEQ:4583
                                                                                                                                                                                                                                                                                                                                                                                                                      09-NOV-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                AAI77642;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAI77642 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          proteins,
                                                                therapy
                                                                          Polymorphic nucleic acid sequences, useful
                                                                                                       WPI; 2001-356160/37
                                                                                                                                Shimkets RA,
                                                                                                                                                                                      30-NOV-1999;
29-NOV-2000;
                                                                                                                                                                                                                             30-NOV-2000; 2000WO-US32758
                                                                                                                                                                                                                                                                                  WO200140521-A2
                                                                                                                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                           (CURA-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1057 CARACCTGATCACACATCAGAGGACACACTCAGGGGGAGAAACCTTATG 1104
                                                                                                                                                                                                                                                                                                                                                                 single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ب
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               . Similarity 81.; 39; Conservative
                                                                                                                                                           CURAGEN CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CAAACCTGATTGTACATCAGAGAACTCATACAGGAGAGAAACCCTATG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           , cytochromes, kinesins, cytokines, interferons,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              invention relates to oligonucleotides encoding polymorphic
                                                                                                                                                                                      2000US-0726173
                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                  py; vaccine; probe; diagnostic assay;
restorative therapy; polymorphic; ds.
                                                                                                                                Leach
                                                                                                                                                                                                    99US-0168138
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Α.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1.2%;
81.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   11 c;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ВР
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  10 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 33.6
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   11 T; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             3.6;
2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     other;
                                                                             ij
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               9;
                                                                           genetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 cyclins, polymerases, treating, e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length
                                                                                                                                                                                                                                                                                                                                                   detection;
                                                                                                                                                                                                                                                                                                                                                              gene therapy;
                                                                           testing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          51;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           interleukins,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           48
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                                                                           and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
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AAI73060 to AAI79867 represent isolated human polymorphic polynucleotide sequences (I), which contain single nucleotide polymorphisms (SNPs).

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RESULT 8
AAL26920
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CC AAM53114 to AAM53329 represent peptides related to human polymorphic CC polynucleotide sequences. The sequences can be used in gene and protein CC therapy, and in vaccine production. (I) and the polypeptides encoded by CC them may be used in the prevention, diagnosis and treatment of diseases CC associated with inappropriate expression of polymorphic polypeptides. CC For example, (I) may be used to treat disorders by rectifying mutations CC or deletions in a patient's genome that affect the activity of CC polypeptides by expressing inactive proteins or to supplement the CC patients own production of polypeptide. Additionally, (I) and its CC complementary sequences may also be used as DNA probes in diagnostic CC assays to detect and quantitate the presence of similar nucleic acids in samples, and therefore which patients may be used as antigens in the CC production of antibodies specific for polymorphic polypeptides. The production of antibodies specific for polymorphic polypeptides. The CC antibodies may also be used as diagnostic agents for detecting the CC The antibodies may also be used as diagnostic agents for detecting the CC presence of polymorphic polypeptides in samples.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local
                                                                                   Polymorphic nucleic acids encoding e.g. amylases, cyclins, polymerases, oncogenes and histones, useful for diagnosing and treating, e.g. cancer, autoimmune diseases and infections -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Immunosuppressive; immunostimulatory; antiinflammatory; cytostatic; neuroprotective; antimicrobial; gene therapy; vaccine; amylase; cancer; amyloid protein; angiopoietin; apoptosis related protein; cadherin; cytoln; polymerase; oncogene; histone; kinase; colony stimulating factor; complement related protein; cytochrome; kinesin; cytokine; interferon; interleukin; G-protein coupled receptor; thioesterase; inflammation;
                                                  Claim 1;
                                                                                                                                                                                          Shimkets RA,
                                                                                                                                                                                                                                                           28-DEC-1999; 99US-0173419
27-DEC-2000; 2000US-0173419
                                                                                                                                                                                                                                                                                                                                                   05-JUL-2001
                                                                                                                                                                                                                                                                                                                                                                                    WO200147944-A2
                                                                                                                                                                                                                                                                                                               28-DEC-2000; 2000WO-US35498
                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAL26920 standard; DNA; 51
                                                                                                                                                                                                                          (CURA-) CURAGEN CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           multifactorial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human SNP oligonucleotide #128
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence
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                                               Page 1434; 4143pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AGTCAAATGCCCAGGCAGATAGGGGTGGGTACCTGGTGAAACCCAACCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AGACAAATGCCTAGGCAGATAGGGGCAGGTCAACAGTGAAACCCCCACCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ВP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                          Leach M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                       disease; autoimmune disease; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A; 15 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1.2%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 33;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             છ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     disease; infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         22;
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The present variants of

relates to related to a

o oligonucleotides amylases, amyloid

proteins,

polymorphic, angiopoietin

the

corresponding

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RESULT 9
ABN33655
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         apoptosis related proteins, cadherin, cyclin, polymerase, oncogenes, histones, kinases, colony stimulating factors, complement related proteins, cytochromes, kinesins, cytokines, interferons, interleukins, G-protein coupled receptors and thioesterases. The present sequence is one such oligonucleotide. The oligonucleotides and the peptides encoded by them may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate expression of the proteins listed above. Disorders that may be prevented, diagnosed and/or treated include multifactorial diseases with a genetic component, such as autoimmune diseases (e.g. rheumatoid arthritis, multiple sclerosis, diabetes, systemic lupus erythromatosus and Grave's disease), inflammation, cancer (e.g. cancers of the bladder, brain, breast, colon and kidney, longariems.
                                                                       The present invention describes oligonucleotide libraries for determines and the populate a (sub-)transcriptome, where the (sub-)transcriptome comprises messenger RNAs transcribed from mult transcription units that populate a genome. The library comprises several oligonucleotides, each capable of hybridising selectively
                                            set of messenger RNAs transcribed from a given transcription unit the genome, which encodes one or more messenger RNA splice variant
          biological
                              The
                                                                                                                                                                          Example 1; SEQ ID 6403; 47pp; English.
                                                                                                                                                                                                           developmental-specific
                                                                                                                                                                                                                      New oligonucleotide libraries comprising oligonucleotides which selectively hybridize to mRNAs transcribed from a transcription a genome, useful for detecting tissue-, pathology-, and
                                                                                                                                                                                                                                                                                                                            Shoshan
                                                                                                                                                                                                                                                                                                                                                                                         28-JUL-2000;
02-MAY-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human spliced transcript detection oligonucleotide SEQ ID NO:6403
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABN33655 standard; DNA; 60
                                                                                                                                                                                                                                                                                                                                                          (COMP-) COMPUGEN INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO200210449-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       splice
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local
           genome, which encodes one or more messenger RNA splice variants. oligonucleotide libraries are useful for detecting mRNAs from a logical sample, in expression profiling studies, in qualitatively
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1412 AAGGACACACAGGAGAGAGCCTTATGTATGCACAGAATGTGGG 1457
                                                                                                                                                                                                                                                                                          2002-257383/30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       mouse; ra
variant;
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37; Conserv
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                                                                                                                                                                                                                                                                                                                         Wasserman
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BP; 19
                                                                                                                                                                                                                                                                                                                                                                                         2000US-221607P
2001US-287724P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     rat;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    transcriptome; oligonucleotide library;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     splice transcript; detection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Α.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                8 C;
                                                                                                                                                                                                                                                                                                                         Α,
                                                                                                                                                                                                           genes
                                                                                                                                                                                                                                                                                                                         Mintz
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               17 G; 7 T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0,:
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Pred. No. 12;
0; Mismatches
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                                                                                                                                                                                                                                                                                                                         Mintz
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0 other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                22;
                                                                                                                                                                                                                                                                                                                       Faigler
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                9;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RNA transcript;
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                                                                                                          from multiple
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                                                                                                                                         detecting
                                                                                                                                                                                                                                           unit of
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RESULT 10
ABN34117
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                 New oligonucleotide libraries comprising oligonucleotides which selectively hybridize to mRNAs transcribed from a transcription a genome, useful for detecting tissue-, pathology-, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        the exemplification of the present invention.

N.B. The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                only expressed in specific tissue under a specific pathological condition; to detect developmental specific genes; and to detect RNA transcripts and splice variants of a transcriptome of a patient suffering
the genome, which encodes one or more messenger RNA splice variants. The oligonucleotide libraries are useful for detecting mRNAs from a biological sample, in expression profiling studies, in qualitatively quantitatively characterising the corresponding transcriptome, and in
                                                                                                            several oligonucleotides, each capable of hybridising selectively to set of messenger RNAs transcribed from a given transcription unit of
                                                                                                                                                               The present invention describes oligonucleotide libraries for detecting messenger RNAs that populate a (sub-)transcriptome, where the (sub-)transcriptome comprises messenger RNAs transcribed from multiple transcription units that populate a genome. The library comprises
                                                                                                                                                                                                                                                                                                                                                             developmental-specific genes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2002-257383/30.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     28-JUL-2000;
02-MAY-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             from a particular disorder. ABN27253 to ABN59589 represent oligonucleotide sequences from rats, humans and mice, which
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                particular biological or pathological state, and so allowing the detection of tissue- and pathology-specific genes such as those genes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   detecting RNA transcripts and splice variants of human or animal transcriptomes. The libraries may also be used as specialised mini libraries to detect transcripts of a sub-transcriptome under a
                                                                                                                                                                                                                                                                                                          Example 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (COMP-) COMPUGEN INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20-JUL-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1419 CACACAGGAGAGAGCCTTATGTATGCACAGAATGTGGG 1457
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          mouse; ra
variant;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       l Similarity
34; Conser
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                                                                                                                                                                                                                                                                                                    SEQ ID 6865; 47pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Wasserman
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
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2001US-287724P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      transcript detection oligonucleotide SEQ ID NO:6865
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          transcriptome;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 splice transcript; detection; RNA tr
anscriptome; oligonucleotide library;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mintz E,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ВΡ
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0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mintz L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB
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messenger RNAs that populate a (sub-)transcriptome, where the (sub-)transcriptome comprises messenger RNAs transcribed from multiple transcription units that populate a genome. The library comprises several oligonucleotides, each capable of hybridising selectively to a set of messenger RNAs transcribed from a given transcription unit of

present invention describes oligonucleotide libraries

for

detecting

biological sample, in expression profiling studies, in qualita quantitatively characterising the corresponding transcriptome,

genome, which encodes one or more messenger RNA splice variants. oligonucleotide libraries are useful for detecting mRNAs from a logical sample, in expression profiling studies, in qualitatively or

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Best Local
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                                                                                                                                                                                       New oligonucleotide libraries comprising oligonucleotides which selectively hybridize to mRNAs transcribed from a transcription a genome, useful for detecting tissue-, pathology-, and
                                                                                                                                                                                                                                                                                                                                  28-JUL-2000;
02-MAY-2001;
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                                                                                                                                                                             developmental-specific genes
                                                                                                                                                                                                                                                   WPI; 2002-257383/30
                                                                                                                                                                                                                                                                             Shoshan A,
                                                                                                                                                                                                                                                                                                                                                                            20-JUL-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human spliced transcript detection oligonucleotide SEQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             15-JUL-2002
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                                                                                                                                                Example 1; SEQ ID 6992; 47pp; English
                                                                                                                                                                                                                                                                                                       (COMP-) COMPUGEN INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ftp.wipo.int/pub/published_pct_sequences.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      mouse; rat; splice transcript;
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39; Conserv
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                                                                                                                                                                                                                                                                           Wasserman
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2001US-287724P
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73.6%;
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oligonucleotide library; ss.
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                                                                                                                                                                                                                                                                           Mintz L,
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Best Local S
Matches 41
                                    messenger RNAs that populate a (sub-)transcriptome, where the (sub-)transcriptome comprises messenger RNAs transcribed from multiple transcription units that populate a genome. The library comprises several oligonucleotides, each capable of hybridising selectively to a set of messenger RNAs transcribed from a given transcription unit of the genome, which encodes one or more messenger RNAs plice variants. The oligonucleotide libraries are useful for detecting mRNAs from a The oligonucleotide libraries are useful for detecting mRNAs from a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           only expressed in specific tissue under a specific pathological condition; to detect developmental specific genes; and to detect RNA transcripts and splice variants of a transcriptome of a patient suffering from a particular disorder. ABN27253 to ABN9589 represent oligonucleotide sequences from rats, humans and mice, which are used in the exemplification of the present invention.

N.B. The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO
                                                                                                                                                                                                                                                                                                                                                                     New oligonucleotide libraries comprising oligonucleotides which selectively hybridize to mRNAs transcribed from a transcription a genome, useful for detecting tissue-, pathology-, and
                                                                                                                                                                                                                                                                                                  Example
                                                                                                                                                                                                                                                                                                                                                       developmental-specific
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02-MAY-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human spliced
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                15-JUL-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (COMP-)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2002-257383/30
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                                                                                                                                                                                                                                                                                                  SEQ ID
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BP;
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2001US-287724P
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                                                                                                                                                                                                                                                                                         7153; 47pp; English.
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anscriptome; oligonucleotide library; ss.
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0; Mismatches
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58;
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biological sample,

characterising

in expression profiling studies, in qualita racterising the corresponding transcriptome,

in qualitatively or

(sub-)transcriptome comprises messenger RNAs transcribed from multiple transcription units that populate a genome. The library comprises several oligonucleotides, each capable of hybridising selectively to a set of messenger RNAs transcribed from a given transcription unit of the genome, which encodes one or more messenger RNA splice variants. The oligonucleotide libraries are useful for detecting mRNAs from a biological sample, in expression profiling studies, in qualitatively or

profiling studies, in qualitatively corresponding transcriptome, and in

or

Example 1; SEQ ID 20382; 47pp; English.

messenger RNAs that

present invention describes oligonucleotide libraries senger RNAs that populate a (sub-)transcriptome, where

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Best Local Similarity
Matches 40; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         detecting RNA transcripts and splice variants of human or animal transcriptomes. The libraries may also be used as specialised mini libraries to detect transcripts of a sub-transcriptome under a particular biological or pathological state, and so allowing the detection of tissue- and pathology-specific genes such as those genes only expressed in specific tissue under a specific pathological condition; to detect developmental specific genes; and to detect RNA transcripts and splice variants of a transcriptome of a patient suffering from a particular disorder. ABN27253 to ABN59589 represent oligonucleotide sequences from rats, humans and mice, which are used in the exemplification of the present invention.

N.B. The sequence data for this patent did not form part of the printed aspecification, but was obtained in electronic format directly from WIPO
                                                                                                                                                                              New oligonucleotide libraries comprising oligonucleotides which selectively hybridize to mRNAs transcribed from a transcription a genome, useful for detecting tissue-, pathology-, and
                                                                                                                                                                                                                                                                                                                    28-JUL-2000;
02-MAY-2001;
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                                                                                                                                                                    developmental-specific genes
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human spliced
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABN47634 standard;
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variant; transcriptome; oligon
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                                                                                                                                                                                                                                                                                          COMPUGEN INC
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79453
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N.B. The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO are format directly from WIPO are the property of the printed are consistent of the printed 
polynucleotide sequences. The sequences can be used in gene and protein therapy, and in vaccine production. (I) and the polypeptides encoded by them may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate expression of polymorphic polypeptides. For example, (I) may be used to treat disorders by rectifying mutations or deletions in a patient's genome that affect the activity of polypeptides by expressing inactive proteins or to supplement the patients own production of polypeptide. Additionally, (I) and its
                                                                                                                                                                                                                                                                                                                                                                           AAI73060 to AAI79867 represent isolated human polymorphic polynucleotide sequences (I), which contain single nucleotide polymorphisms (SNPs). AAM53114 to AAM53329 represent peptides related to human polymorphic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2001-356160/37
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 1; Page 2464; 2653pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Polymorphic nucleic acid sequences, useful in genetic testing
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        polynucleotide sequences. The sequences can be used in gene and protein therapy, and in vaccine production. (I) and the polypeptides encoded by them may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate expression of polymorphic polypeptides. For example, (I) may be used to treat disorders by rectifying mutations or deletions in a patient's genome that affect the activity of polypeptides by expressing inactive proteins or to supplement the patients own production of polypeptide. Additionally, (I) and its complementary sequences may also be used as DNA probes in diagnostic assays to detect and quantitate the presence of similar nucleic acids in samples, and therefore which patients may be in need of restorative therapy. The polypeptides encoded by (I) may be used as antigens in the production of antibodies specific for polymorphic polypeptides. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      complementary sequences may also be used as DNA probes in diagnostic assays to detect and quantitate the presence of similar nucleic acids in samples, and therefore which patients may be in need of restorative therapy. The polypeptides encoded by (I) may be used as antigens in the production of antibodies specific for polymorphic polypeptides. The antibodies may also be used to down regulate expression and activity. The antibodies may also be used as diagnostic agents for detecting the presence of polymorphic polypeptides in samples.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; single nucleotide polymorphism; SNP; genome; protein therapy; vaccine; probe; diagnostic assay; c quantitation; restorative therapy; polymorphic; ds.
                                                                                                                                                                                                                               AA173060 to AA179867 represent isolated human polymorphic polynucleotide sequences (I), which contain single nucleotide polymorphisms (SNPs). AAM53114 to AAM53329 represent peptides related to human polymorphic
                                                                                                                                                                                                                                                                                                    Claim 1; Page 2050; 2653pp; English
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                                                                                                Matches
                                                                                                                   Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local
                                                                                                                                                                                        Sequence
                                                                                                                                                                                                                                                 virus), modulating gene expression, detecting an altered zinc finger recognition sequence and diagnosing disease associated with abnormal genomic structure. The present DNA sequence is zinc finger domain 1 (Zif-1) specific PCR primer 1 used to generate three finger ZFP. This
                                                                                                                                                                                                                                                                                                                                 the identification of a context-independent recognition code to design zinc finger domains. The invention also relates to zinc finger proteins (ZFP) designed using this recognition code. The ZFPs are useful for altering genomic structure, inhibiting viral replication (where viral replication is inhibited for plant virus, an animal virus or a human replication is inhibited for plant virus, an animal virus or a human replication is inhibited for plant virus, an animal virus or a human replication is inhibited for plant virus, an animal virus or a human virus or a
                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present invention relates to novel DNA binding proteins comprising zinc finger domains in which two histidine and two cysteine residues coordinate a central zinc ion. The invention particularly relates to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DNA binding protein; zinc finger domain 1; zinc finger protein; viral replication; gene expression; virucide; Zifl; PCR; primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2002-172000/22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     31-JAN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The antibodies may also presence of polymorphic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   for diagnosing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New Zinc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (SYGN ) SYNGENTA PARTICIPATIONS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         21-JUL-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      19-JUL-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO200208286-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Zif-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    01-JUL-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAD33614;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAD33614 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 51 BP;
                                                    1593
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1687 CATTTGTATGTGCTGAGTGTGGACGAGGCTTTAATGATAAGTCCACCCTC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     specific PCR primer 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             5
                                                                                            Similarity
36; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  36; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      6;
                          GGGGAGAAGCCATTTGTATGTACGGAGTGTGGGCGAGGCTTTACCCGGGAAATCAACCCTG
                                                                                                                                                                                        60
GGGGAGAAGCCGTATAAATGTCCGGAATGTGGTAAAAGTTTTAGCNNNAGCNNNNNNTTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Finger Protein (ZFP) comprising three essential nosing diseases associated with abnormal genomic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CTTATGAATGTACAGAGTGTGGGAAAGCCTTCAAACATAGTTCCACCCTC 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Page 70; 143pp; English.
                                                                                                                                                                                      ВP;
                                                                                                                                                                                                                                     used
                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2000US-220060P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2001WO-EP08367
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                      15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          13
                                                                                                                                                                                                                              in the exemplification of the
                                                                                                                                                                                   A; 6 C; 17 G; 13 T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A; 10 C;
                                                                                                            1.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       72.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  be used as diagnostic agents polypeptides in samples.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ВP
                                                                                            0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                used to generate three
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     13 G; 15 T; 0 other;
                                                                                                               Score 27;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Pred. No. 2.2e
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score
                                                                                            Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ΑG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     27.6;
No. 2
                                                                           DB 24,
3.8e+02;
24;
                                                                                                                                                                                   9 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       .2e+02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 22;
                                                                                                                                                                                                                                invention
                                                                                                                                  Length 60
                                                                                         Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  finger ZFP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       for detecting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              domains useful structure -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          51;
                                                                                         0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1736
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0;
                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ZFP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         the
  6.0
                                            ..652
                                                                                       0;
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AAL33199 ID AAL3

AAL33199 standard; DNA; 51 BP

RESULT 18

망 Q

Matches Query Match

34;

Conservative

0;

Pred. No. 4e+ 0; Mismatches Score 26.8; Pred. No. 46

4e+02;

Indels Length

0

Gaps

0

259

DB 22; 12;

51;

214 TGGCATTCAGGGATGTGGCTGTGTACTTCACCCAGGAGGAGTGGAG

TGACTTTTGAAGACGTGGCTGTGAAATTTTCCCAGGAGGAATGGAG

Local

Similarity

1.0%; 73.9%;

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RESULT 17
AAL32346
                                                                                                                                                                                                                                                                                                          G-protein coupled receptors and thioesterases. The present sequence is one such ollyonucleotide. The ollyonucleotides and the peptides encoded by them may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate expression of the proteins listed above. Disorders that may be prevented, diagnosed and/or treated include multifactorial diseases with a genetic component, such as autoimmune diseases (e.g. rheumatoid arthritis, multiple sclerosis, diabetes,
                                                                                                                                               systemic lupus erythromatosus and Grave's disease), inflammation, (e.g. cancers of the bladder, brain, breast, colon and kidney, leukaemia), diseases of the nervous system and an infection of particular color of the partic
           Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       variants of proteins related to amylases, amyloid proteins, angiopoiet apoptosis related proteins, cadherin, cyclin, polymerase, oncogenes, histones, kinases, colony stimulating factors, complement related proteins, cytochromes, kinesins, cytokines, interferons, interleukins,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Polymorphic nucleic acids encoding e.g. amylases, oncogenes and histones, useful for diagnosing and cancer, autoimmune diseases and infections -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 1; Page 2986; 4143pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Shimkets
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          28-DEC-1999;
27-DEC-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         28-DEC-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               complement related protein; cytochrome; kinesin; cytokine; interfero interleukin; G-protein coupled receptor; thioesterase; inflammation; multifactorial disease; autoimmune disease; infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          amyloid protein; angiopoietin; apoptosis related protein; cadherin; cyclin; polymerase; oncogene; histone; kinase; colony stimulating factor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAL32346 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        05-JUL-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        nervous system disease; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              neuroprotective;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human SNP oligonucleotide #5554.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (CURA-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Immunosuppressive;
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           51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RΑ,
           BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             invention relates to oligonucleotides encoding polymorphic proteins related to amylases, amyloid proteins, angiopoietin,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    99US-0173419
2000US-0173419
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Leach
           12
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antimicrobial; gene therapy; vaccine; amylase; can
           Α.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       3
           9
           Ç
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           15 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ВP
     15 T; 0
other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              cyclins, polymerases,
treating, e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   interferon;
                                                                                                                                                          pathogenic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              cancer;
                                                                                                                                                                                                                                                                    cancer
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RESULT 19
ABL56734
ID ABL56
XX
                                                                                                                     QΥ
                                                                                                                                                                                                 XX
AC
                                                                                               В
                                                                                                                                                          Query Match
Best Local :
                                                                                                                                                Matches
                                                                                                                                                                                                                                                                               apoptosis related proteins, cadherin, cyclin, polymerase, oncogenes, histones, kinases, colony stimulating factors, complement related proteins, cytochromes, kinesins, cytokines, interferons, interleukins, G-protein coupled receptors and thioesterases. The present sequence is one such oligonucleotide. The oligonucleotides and the peptides encoded by them may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate expression of the proteins listed above. Disorders that may be prevented, diagnosed and/or treated include multifactorial diseases with a genetic component, such as autoimmune
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Polymorphic nucleic acids encoding e.g. amylases, cyclins, polymerases, oncogenes and histones, useful for diagnosing and treating, e.g. cancer, autoimmune diseases and infections -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  complement related protein; cytochrome; kinesin; cytokine; interferon; interleukin; G-protein coupled receptor; thioesterase; inflammation; multifactorial disease; autoimmune disease; infection; nervous system disease; ss.
         ABL56734;
                                                                                                                                                                                                                                                                                                                                                                                                 The present invention relates to oligonucleotides encoding polymorphic variants of proteins related to amylases, amyloid proteins, angiopoietin,
                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAL33199
                                   ABL56734 standard;
                                                                                                                                                                                                Sequence
                                                                                                                                                                                                                                                               systemic
                                                                                                                                                                                                                                                                          diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2001-465210/50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Shimkets
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           28-DEC-1999; 99US-0173419
27-DEC-2000; 2000US-0173419
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                28-DEC-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          05-JUL-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO200147944-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cyclin;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           neuroprotective;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Immunosuppressive; immunostimulatory; antiinflammatory; cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          24-JAN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (CURA-) CURAGEN CORP
                                                                                                                                                           Local Similarity
                                                                                                                                                                                                                                    seases (e.g. rheumatoid arthritis, multiple scierosis, diabetes, stemic lupus erythromatosus and Grave's disease), inflammation, cancer.g. cancers of the bladder, brain, breast, colon and kidney, ukaemia), diseases of the nervous system and an infection of pathogeni
                                                                                                                     1560 TCAAACCTTAACAAACACCAGAGGTCACACGGGGGGGAGAAGCCAT 1605
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SNP oligonucleotide #6407
                                                                                               Н
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                protein; angropoletin; apoptosis related protein; cadherin; polymerase; oncogene; histone; kinase; colony stimulating factor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RA,
                                                                                                                                                                                                 51
                                                                                                                                                                                                                                                                                                                                                                                                                                       Page 3217; 4143pp; English.
                                                                                               TCAAACCTGCACATCCACCAGCGAGAACACACAGGAGAGAGGCCCCT
                                                                                                                                                                                                BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2000WO-US35498
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Leach M;
                                                                                                                                                                                                 19
                                                                                                                                                                                                A; 17
                                   DNA;
                                                                                                                                                          1.0%;
                                   41
                                                                                                                                                                                               C;
                                   ВР
                                                                                                                                                                                               11 G; 4 T; 0 other;
                                                                                                                                                0;
                                                                                                                                             Score 26.8; I
Pred. No. 4e+(
0; Mismatches
                                                                                                                                                         8; DB 22;
4e+02;
                                                                                                                                              12;
                                                                                                                                                                     Length 51;
                                                                                                                                              Indels .
                                                                                               46
                                                                                                                                             0;
                                                                                                                                                                                                                                    pathogenic
                                                                                                                                             Gaps
                                                                                                                                              0,
                                                                                                                                                                                                                                                                                                                                                                                                                            В
                                                                                                                                                                                                                                                                                                                                                  Q
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RESULT 20
AAS18906
ID AAS18
XX AAS18
XX AAS18
XX Humar
XX Humar
XX Humar
XX Humar
XX Homo
COS Homo
COS Homo
COS Homo
COS 103-OC
XX CN131
X
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Probes ABL56733-34 were used to identify cDNA encoding a human krupple zinc finger related protein 70.84. The polypeptide and encoding polynucleotide are used in diagnosis and treatment of
                                                                                                                                                                                                                                                              CN1315334-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              malignant tumour, haemal disorders, human immunodeficiency virus infection, immunological diseases and various inflammations. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human Krupple Zinc Finger related protein 70.84 and encoding polynucleotide, useful in the diagnosis and treatment of malignant tumours, haemal disorders, human immunodeficiency virus infection,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mao
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     04-APR-2002
                                                                 27-MAR-2000;
                                                                                                                                  27-MAR-2000;
                                                                                                                                                                                                   03-OCT-2001.
                                                                                                                                                                                                                                                                                                                            Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human OZF gene related protein 13 oligonucleotide probe #1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAS18906;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAS18906 standard; DNA; 41
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               polynucleotide may also be used for gene therapy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 7; Page 15; 39pp; Chinese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (BIOW-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  07-JUL-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   02-JUL-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WO200226807-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  gene therapy; probe;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        naemar
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      30-JUL-2002
(BODE-) BODE GENE DEV CO LTD SHANGHAI
                                                                                                                                                                                                                                                                                                                                                                                                 human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  26-MAR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                immunological diseases and inflammation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2002-292482/33.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 301
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    #2
                                                                                                                                                                                                                                                                                                                                                                                             ; ss; OZF gene related p
immunodeficiency virus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 krupple zinc finger related protein 70.84; tumour; disorder; HIV infection; immunological disease; inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   <u>--</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  l Similarity
32; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BIOWINDOW
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Xie Y;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     41
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BP; 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2000CN-0117012.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    krupple zinc finger related protein 70.84
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                 2000CN-0115166
                                                                                                                                  2000CN-0115166
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A; 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENE DEV INC SHANGHAI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1.0%;
78.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 C; 12 G; 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ВÞ
                                                                                                                                                                                                                                                                                                                                                                                             protein 13;
s; probe.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 26.6; DB 24;
Pred. No. 4e+02;
D; Mismatches 9;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                            cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                               VIΗ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length
                                                                                                                                                                                                                                                                                                                                                                                                                                   infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   41;
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RESULT 21
AAHE9605/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local S
Matches 32
     glomerulonephritis, haemolytic
          aberrant protein expression, including cancer, amyloidosis Alzheimer's disease, Down's syndrome, oedema, lupus (SLE),
                                    The present invention relates to human nucleic acids containing single nucleotide polymorphisms (SNPs). These can be used in forensic and paternity tests, and to aid in the treatment of diseases associated with
                                                               The present invention relates to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to a new polypeptide-OZF (not defined) gene related protein 13, the polypnucleotide encoding it, the process for preparing the said polypeptide by DNA recombination, the application of said polypeptide in treating several diseases such as cancer and HIV infection (human immunodeficiency virus). The antagonist against the said polypeptide and its therapeutic action, and the application of said polypeptide and its therapeutic action, and the application of said polypeptide for coding this new-OZF correlated protein 13 are disclosed. The present sequence is -OZF correlated protein to action the present sequence is -OZF correlated protein 13 are
                                                                                              Claim 1;
                                                                                                                    polymorphisms, infection and
                                                                                                                                                                           P-PSDB;
                                                                                                                                                                                                                                                                 07-JAN-2000;
                                                                                                                                                                                                                                                                                          05-JAN-2001; 2001WO-US00322
                                                                                                                                                                                                                                                                                                                                             WO200151670-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                  Human DNA/RNA binding protein DNA polymorphic
                                                                                                                                                                                                                                        (CURA-) CURAGEN CORP
                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                               Human; single forensic test.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                01-OCT-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAH89605;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAH89605
                                                                                                                                                Isolated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 41 BP; 15 A; 6 C; 11 G; 9 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Polypeptide-OZF
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1411 TAAGGACACACACAGGAGAGAAGCCTTATGTATGCACAGAA
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                                                                                                                                                                          2001-451871/48.
DB; AAM00488.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            nucleic
                                                                                                                 human polynucleotides containing single nucleotide hisms, useful for the treatment and diagnosis of e.un and diabetes
                                                                                          Page 213; 475pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Xie Y;
                                                                                                                                                                                                                                                                                                                                                                                                test;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TGAGAAGTCACAGGTGAGAAGCCCTATGTATGTAATGAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                 2000US-0174962
                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                               Leach
                                                                                                                                                                                                                                                                                                                                                                                            nucleotide polymorphism;
aberrant protein express
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         encoding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  51
                                                                                           English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Chinese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ΒP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Pred.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OZF gene related protein 13
  anaemia,
                                                                                                                                                                                                                                                                                                                                                                                            rphism; SNP;
expression;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          4e+02;
thrombocytopaenia,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             and polynucleotide encoding
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                                                                                                                                                                                                                                                                                                                                                                                              paternity test; ds.
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                                                                                                                                                                                                                                                                                                                                                                                                                                   site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               9;
                           amyloidosis, diabetes,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQ
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                                                                                                                                of e.g.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         41;
                                                                                                                                                                                                                                                                                                                                                                                                                                   NO:
             vasculitis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0;
                                                                                                                                cancer,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0;
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δÃ

921 GGGGAGACACCTTACATGTACACTGAGTGGGGAGACAGCTTTGGCAGTA 969

Matches Query Match Best Local

35;

Conservative

0,

Pred. No. 4.9 0; Mismatches Score 26.6; Pred. No. 4

9e+02; DB 24;

Indels Length

0;

Gaps

0

56;

Similarity

1.0%;

Sequence

56

BP;

15 P.

10 C;

19

G;

12 T; 0 other;

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RESULT 22
AAD33627
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         В
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  8×888888
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Qy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Ouery Match
Best Local S
Matches 35
                                                          altering genomic structure, inhibiting viral replication (where viral replication is inhibited for plant virus, an animal virus or a human virus), modulating gene expression, detecting an altered zinc finger recognition sequence and diagnosing disease associated with abnormal genomic structure. The present DNA sequence is zinc finger domain 3 (Zif-3) specific oligonucleotide which is used to generate three finger.
                                                                                                                                                 the identification of a context-independent recognition code to design zinc finger domains. The invention also relates to zinc finger proteins (ZFP) designed using this recognition code. The ZFPs are useful for
                                                                                                                                                                                                 The present invention relates to novel DNA binding proteins comprising zinc finger domains in which two histidine and two cysteine residues zonc finate a central zinc ion. The invention particularly relates to the invention particularly relates to
                                                                                                                                                                                                                                                                 Example 7; Page 72; 143pp;
                                                                                                                                                                                                                                                                                                                                                 WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                           21-JUL-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          19-JUL-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       31-JAN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO200208286-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DNA binding protein; viral replication; ge
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          meningitis, muscular disorders, dementia, neurological diseases, tuberous sclerosis, male infertility, hypercalcaemia, blood pressure disorders, osteoporosis, pathogenic infections, hypercholesterolaemia, obesity or autoimmunity. The present sequence is a polymorphism-containing oligonucleotide fragment of the invention.
                                                                                                                                                                                                                                                                                                     for
                                                                                                                                                                                                                                                                                                                                                                                                            (SYGN ) SYNGENTA PARTICIPATIONS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Zif-3 specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             01-JUL-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAD33627
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAD33627 standard; DNA; 56
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 51
                                               for
                                                                                                                                                                                                                                                                                              Zinc Finger Protein (ZFP) comprising three essential diagnosing diseases associated with abnormal genomic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1960 GCTTTAGCCGGCAGTCACACCTCATTAGACACCAGAGGACACATTCAGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         49
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  35; Conser
                                            specific oligonucleotide which
the L1 site of beet curly top v
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GCTTCAGCCGCCCTCGCACCTGCTGCGCCACCAGCGGACTCACACAGG
                             LI site of beet curly top virus (BCTV). exemplification of the invention.
                                                                                                                                                                                                                   invention relates to novel DNA domains in which two histidine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BP; 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                           2000US-220060P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2001WO-EP08367
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A; 13 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             oligo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 zinc finger domain 3; zir
ene expression; virucide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 71.48;
                                                                                                                                                                                                                                                               English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ВР
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              used
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               G; 9 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                            AG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             to generate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                         sequence is zinc finger domain 3 hich is used to generate three finger
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               26.6;
No. 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               6e+02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               zinc findide; Zif3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DВ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             three finger
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             finger protein;
if3; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length
                                        This sequence
                                                                                                                                                                                                                                                                                                structure
                                                                                                                                                                                                                                                                                                               domains useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                51;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2008
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
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RESULT 23
AAI79452
                               Ωy
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                                                                                                                                                             therapy, and in vaccine production. (I) and the polypeptides encoded by them may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate expression of polymorphic polypeptides. For example, (I) may be used to treat disorders by rectifying mutations or deletions in a patient's genome that affect the activity of polypeptides by expressing inactive proteins or to supplement the patients own production of polypeptide. Additionally, (I) and its complementary sequences may also be used as DNA probes in diagnostic assays to detect and quantitate the presence of similar nucleic acids in samples, and therefore which patients may be in need of restorative therapy. The polypeptides encoded by (I) may be used as antigens in the production of antibodies specific for polymorphic polypeptides. The antibodies may also be used to down regulate expression and activity. The antibodies may also be used as diagnostic agents for detecting the presence of polymorphic polypeptides in samples.
                                                                                 Query Match
Best Local Similarity
                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human; single nucleotide polymorphism; SNP; genome; gene ther protein therapy; vaccine; probe; diagnostic assay; detection; quantitation; restorative therapy; polymorphic; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          07-JUN-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human silent SNP containing nucleic acid SEQ:6393.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             09-NOV-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                            sequences (I), which contain single nucleotide polymorphisms AAM53114 to AAM53329 represent peptides related to human poly
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAI73060 to AAI79867 represent isolated human polymorphic polynucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Polymorphic nucleic acid sequences, useful in genetic testing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2001-356160/37
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     30-NOV-1999;
29-NOV-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         30-NOV-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAI79452 standard;
                                                                                                                                      Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                            polynucleotide sequences. The sequences can be used in gene and protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    tnerapy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (CURA-) CURAGEN CORP.
                                 1243
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1; Page 2464; 2653pp;
 \vdash
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RΑ,
                                                                                                                                  51 BP; 16 A; 7 C; 18 G;
AGAGAGTCCACACTGGAGAGAAACCCTATAGATGTTGTGGATGTGGG
                               AGAGGGCGCACACTGGGGAGAAGCCTTATGTTTGCAGGGAATGTGGG 1289
                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     99US-0168138.
2000US-0726173.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2000WO-US32758.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Leach M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DNA;
                                                                                 0.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ВР
                                                                   0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 English
                                                               Score 26.2; DB 2
Pred. No. 6.2e+02
0; Mismatches 1
                                                                                                                                  10 T; 0 other;
                                                                                                 DB 22;
                                                                   Indels
                                                                                             Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          gene therapy;
 47
                                                                                                                                                                                                                                                                                                                                                                                                                                            polymorphic
                                                               0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                             (SNPs)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  and
                                                                 Gaps
                                                                 0,
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RESULT 24 AAI78092/c

01-JUL-2002 AAD33623;

(first entry)

AAD33623 standard;

DNA;

60 ВP

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RESULT 25
AAD3623
ID AAD7
XX
AC A/
XX
OT
XX
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                                                                                                                                                                                            20
                                                                                                                                                                                                                         Query Match
Best Local S
Matches 35
                                                                                                                                                                                                                                                                                                                                For example, (I) may be used to treat disorders by rectifying mutations or deletions in a patient's genome that affect the activity of polypeptides by expressing inactive proteins or to supplement the patients own production of polypeptide. Additionally, (I) and its complementary sequences may also be used as DNA probes in diagnostic assays to detect and quantitate the presence of similar nucleic acids in samples, and therefore which patients may be in need of restorative therapy. The polypeptides encoded by (I) may be used as antigens in the production of antibodies specific for polymorphic polypeptides. The antibodies may also be used as diagnostic agents for detecting the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       polynucleotide sequences. The sequences can be used in gene and protein therapy, and in vaccine production. (I) and the polypeptides encoded by them may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate expression of polymorphic polypeptides.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    sequences (I), which contain single nucleotide polymorphisms (SNPS).

AAM53114 to AAM53329 represent peptides related to human polymorphic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAI73060 to AAI79867 represent isolated human polymorphic polysequences (I), which contain single nucleotide polymorphisms
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 1; Page 2050; 2653pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              30-NOV-1999;
29-NOV-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; single nucleotide polymorphism; SNP; genome; protein therapy; vaccine; probe; diagnostic assay; quantitation; restorative therapy; polymorphic; ds.
                                                                                                                                                                                                                                                                                      Sequence 51 BP; 13 A; 11 C;
                                                                                                                                                                                                                                                                                                                   presence of polymorphic polypeptides in samples.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Polymorphic nucleic acid sequences, useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2001-356160/37
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              30-NOV-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             07-JUN-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO200140521-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human silent SNP containing nucleic acid SEQ:5033
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      09-NOV-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAI78092 standard; DNA; 51 BP
                                                                                                                                                                                          1687 CATTTGTATGTGCTGAGTGTGGACGAGGCTTTAATGATAAGTCCACCCTC 1736
                                                                                                                                                             51
                                                                                                                                                                                                                           э́5;
                                                                                                                                                                                                                                        Similarity
                                                                                                                                                             CTTATGAATGTACAGAGTGTGGGAAGGCCTTCAAACATAGTTCCACCCTC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RA,
                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              99US-0168138.
2000US-0726173.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Leach M;
                                                                                                                                                                                                                                        0.9%;
                                                                                                                                                                                                                         Score 26; DB
Pred. No. 7.2e
0; Mismatches
                                                                                                                                                                                                                           0,
                                                                                                                                                                                                                                                                                      13 G; 14 T; 0 other;
                                                                                                                                                                                                                                                        DB 22;
                                                                                                                                                                                                                                          .2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       in
                                                                                                                                                                                                                           15:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     genetic testing
                                                                                                                                                                                                                                                       Length
                                                                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       detection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    gene therapy;
                                                                                                                                                                                                                                                          51;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        polynucleotide
                                                                                                                                                             Ν
                                                                                                                                                                                                                           0;
                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                           0;
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RESULT 26
AAD33618
ID AAD33
DЬ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local !
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present invention relates to novel DNA binding proteins comprising zinc finger domains in which two histidine and two cysteine residues coordinate a central zinc ion. The invention particularly relates to the identification of a context-independent recognition code to design zinc finger domains. The invention also relates to zinc finger proteins altering genomic structure, inhibiting viral replication (where viral replication is inhibited for plant virus, an animal virus or a human virus), modulating gene expression, detecting an altered zinc finger recognition sequence and diagnosing disease associated with abnormal genomic structure. The present DNA sequence is zinc finger domain 1 (Zif-1) specific oligonucleotide which is used to generate three finger for the related to the sequence is zinc finger domain 1 (Zif-1) specific oligonucleotide which is used to generate three finger
           19-JUL-2001; 2001WO-EP08367
                                                                                                        Unidentified.
                                                                                                                                 DNA binding protein; zinc finger domain 3; zinc finger protein; ZFP; viral replication; gene expression; virucide; Zif3; PCR; primer; ss.
                                                                                                                                                                                2if-3 specific PCR primer 5 used to generate three finger ZFP.
                                                                                                                                                                                                                   01-JUL-2002 (first entry)
                                                                                                                                                                                                                                                                                 AAD33618 standard; DNA; 56 BP
                                             31-JAN-2002
                                                                         WO200208286-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 7; Page 71; 143pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2002-172000/22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New Zinc Finger Protein (ZFP) comprising three essential domains useful for diagnosing diseases associated with abnormal genomic structure
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   21-JUL-2000; 2000US-220060P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  19-JUL-2001; 2001WO-EP08367
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     31-JAN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO200208286-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DNA binding protein; zinc finger domain 1; zinc finger protein; viral replication; gene expression; virucide; Zifl; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Zif-1 specific sense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (SYGN ) SYNGENTA PARTICIPATIONS AG.
                                                                                                                                                                                                                                                                                                                                                                                                                                             Local
                                                                                                                                                                                                                                                                                                                                                                               1593 GGGGAGAAGCCATTTGTATGTACGGAGTGTGGGCGAGGCTTTACC 1637
                                                                                                                                                                                                                                                                                                                                                                 ш
                                                                                                                                                                                                                                                                                                                                                                                                                            l Similarity
33; Conserv
                                                                                                                                                                                                                                                                                                                                                           GGGGAGAAGCCGTATAAATGTCCGGAATGTGGTAAAAGTTTTAGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         60 BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       L1 site of beet curly top virus (BCTV). exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      18 A; 9 C; 19 G; 14 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                          0.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        oligo used to generate three finger ZFP
                                                                                                                                                                                                                                                                                                                                                                                                                         Score 25.8; DI
Pred. No. 9.2e
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                           0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                          9.2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 24;
                                                                                                                                                                                                                                                                                                                                                                                                                           12;
                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length
                                                                                                                                                                                                                                                                                                                                                              45
                                                                                                                                                                                                                                                                                                                                                                                                                                                          60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  sequence is
                                                                                                                                                                                                                                                                                                                                                                                                                           0;
                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ZFP;
                                                                                                                                                                                                                                                                                                                                                                                                                         0;
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         New oligonucleotide libraries comprising oligonucleotides which selectively hybridize to mRNAs transcribed from a transcription a genome, useful for detecting tissue-, pathology-, and developmental-specific genes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present invention relates to novel DNA binding proteins comprising zinc finger domains in which two histidine and two cysteine residues coordinate a central zinc ion. The invention particularly relates to the identification of a context-independent recognition code to design zinc finger domains. The invention also relates to zinc finger proteins (ZFP) designed using this recognition code. The ZFPs are useful for altering genomic structure, inhibiting viral replication (where viral replication is inhibited for plant virus, an animal virus or a human virus), modulating gene expression, detecting an altered zinc finger recognition sequence and diagnosing disease associated with abnormal genomic structure. The present DNA sequence is zinc finger domain 3 (Zif-3) specific PCR primer 5 used to generate three finger ZFP. This sequence is used in the exemplification of the invention.
                                                                                                               Shoshan A, Wasserman
                                                                                                                                                                         28-JUL-2000;
02-MAY-2001;
                                                                                                                                                                                                                    20-JUL-2001; 2001WO-IB01903
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                                                                                                                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                                                                                                                     Human; mouse; rat; splice transcript; detection; RNA tr splice variant; transcriptome; oligonucleotide library;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABN34081 standard; DNA; 60 BP
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                                                                                                                                            (COMP-) COMPUGEN INC
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                                                                                                                                                                                                                                                                                                                                                                                                                15-JUL-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                             ABN34081;
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31; Conser
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2001US-287724P
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                                                                                                               A,
                                                                                                                 Mintz
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Pred. No. 1e
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Example 1; SEQ ID

6829; 47pp; English.

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RESULT 28
ABN34388
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Best Local S
Matches 37
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          several oligonucleotides, each capable of hybridising selectively to a set of messenger RNAs transcribed from a given transcription unit of the genome, which encodes one or more messenger RNAs splice variants. The oligonucleotide libraries are useful for detecting mRNAs from a biological sample, in expression profiling studies, in qualitatively or quantitatively characterising the corresponding transcriptome, and in detecting RNA transcripts and splice variants of human or animal transcriptomes. The libraries may also be used as specialised mini
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention describes oligonucleotide libraries for detecting messenger RNAs that populate a (sub-)transcriptome, where the (sub-)transcriptome comprises messenger RNAs transcribed from multiple transcription units that populate a genome. The library comprises several oligonucleotides, each capable of hybridising selectively to a
                              New oligonucleotide libraries comprising oligonucleotides which selectively hybridize to mRNAs transcribed from a transcription a genome, useful for detecting tissue-, pathology-, and
                                                                                                                                                                                                                                                                                                                                                                                                                   Human; mouse; ra
splice variant;
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02-MAY-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           N.B. The sequence data for this patent did not form part of the specification, but was obtained in electronic format directly fr
                developmental-specific genes
                                                                                                                                           Shoshan A,
                                                                                                                                                                                                                                                                        20-JUL-2001; 2001WO-IB01903
                                                                                                                                                                                                                                                                                                            07-FEB-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human spliced transcript detection oligonucleotide SEQ ID NO:7136
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                                                                                                                                                                               (COMP-) COMPUGEN INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ftp.wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1190 GTGCAAGGAATGTGGGCAGAGCTTTAGCCTGAAGTCAAACCTCATTACCCACCAGA 1245
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37; Conserv
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                                                                                                                                           Wasserman A,
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t; transcriptome; oligonucleotide library; ss.
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Pred. No. 1.1e
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                                                                                                                                           Mintz L,
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N.B. The sequence data for this patent did not form part of the printed as pecification, but was obtained in electronic format directly from WIPO and the property specification of the present approach to the presence of the printed as pecification, but was obtained in electronic format directly from WIPO and the property specification of the present approach to the present approach to the property specification of the present approach to the 
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                                                                                                                                Polypeptide-human zinc finger and PHD finger structural doman-contg actin 15 and polynucleotide for coding said polypeptide -
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 25.4;
Pred. No. 1.
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.2e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            16;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      disease;
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Example 7; Page 21 (disclosure); 35pp; Chinese

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RESULT. 30
AAL27311
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Best Local :
      proteins, cytochromes, kinesins, cytokines, interferons, interleukins, G-protein coupled receptors and thioesterases. The present sequence is one such oligonucleotide. The oligonucleotides and the peptides encoded by them may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate expression of the proteins listed above. Disorders that may be prevented, diagnosed and/or treated include multifactorial diseases with a genetic component, such as autoimmune diseases (e.g. rheumatoid arthritis, multiple sclerosis, diabetes, systemic lupus erythromatosus and Grave's disease), inflammation, cancer systemic lupus erythromatosus and Grave's disease).
                                                                                                                                                variants of proteins related to amylases, amyloid proteins, angiopoi apoptosis related proteins, cadherin, cyclin, polymerase, oncogenes, histones, kinases, colony stimulating factors, complement related
                                                                                                                                                                            The present invention relates to oligonuc variants of proteins related to amylases.
                                                                                                                                                                                                                                                       Polymorphic nucleic acids encoding e.g. amylases, oncogenes and histones, useful for diagnosing and cancer, autoimmune diseases and infections -
                                                                                                                                                                                                                               Claim
    e
                                                                                                                                                                                                                                                                                                                      WPI; 2001-465210/50
                                                                                                                                                                                                                                                                                                                                                    Shimkets
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                                                                                                                                                                                                                                                                                                                                                                                                                           28-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                       28-DEC-2000; 2000WO-US35498
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Immunosuppressive; immunostimulatory; antiinflammatory; coneuroprotective; antimicrobial; gene therapy; vaccine; amanyloid protein; angiopoietin; apoptosis related protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO200147944-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  HOMO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             nervous
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             multifactorial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   interleukin; G-protein coupled receptor; thioesterase;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to a novel polypeptide, human zinc finger of pHD finger structural domain-contig actin 15. It is useful in the treatment of cancer, nosohaemia, HIV infection, immunological distand inflammatory disorders. The present sequence is a probe used to detect polynucleotides encoding the polypeptide of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           complement
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         cyclin;
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   .g. cancers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1239 CACCAGAGGGCGCACACTGGGGAGAAGCCTTATGTTTG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        30; Conser
                                                                                                                                                                                                                                                                                                                                                                                 CURAGEN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   polymerase; oncogene; histone; kinase; colony stimulating ant related protein; cytochrome; kinesin; cytokine; interf
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             system
                                                                                                                                                                                                                            Page 1538; 4143pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
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of
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                                                                                                                                                                                                                                                                                                                                                                                                                          99US-0173419
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          disease;
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0; Mismatches
 brain,
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Pred. No. 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     immunological diseases
                                                                                                                                                                              encoding polymorphic
proteins, angiopoietin,
                                                                                                                                                                                                                                                                   cyclins, polymerases treating, e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      inflammation;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    interferon;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  factor;
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RESULT
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Best Local
                                        The Oilgonucleotide libraries are useful for detecting mRNAs from a biological sample, in expression profiling studies, in qualitatively or quantitatively characterising the corresponding transcriptome, and in detecting RNA transcripts and splice variants of human or animal transcriptomes. The libraries may also be used as specialised mini libraries to detect transcripts of a sub-transcriptome under a particular biological or pathological state, and so allowing the detection of tissue- and pathology-specific genes such as those genes only expressed in specific tissue under a specific pathological condition; to detect developmental specific genes such as those genes conly expressed in specific tissue under a specific pathological condition; to detect developmental specific genes; and to detect RNA transcripts and splice variants of a transcriptome of a patient suffering from a particular disorder. ABN27253 to ABN59589 represent
               from a particular disorder. ABN27253 to ABN59589 oligonucleotide sequences from rats, humans and the exemplification of the present invention.
                                                                                                                                                                                                                             (sub-)transcriptome comprises messenger RNAs transcribed from multiple transcription units that populate a genome. The library comprises several oligonucleotides, each capable of hybridising selectively to a set of messenger RNAs transcribed from a given transcription unit of
                                                                                                                                                                                                                                                                                       messenger RNAs that populate a (sub-)transcriptome, where
                                                                                                                                                                                                                                                                                                                                                                           New oligonucleotide libraries comprising oligonucleotides which selectively hybridize to mRNAs transcribed from a transcription a genome, useful for detecting tissue, pathology, and
                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI;
                                                                                                                                                                                                                                                                                                                                         Example 1;
                                                                                                                                                                                                                                                                                                                                                                  developmental-specific genes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               28-JUL-2000;
02-MAY-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (COMP-) COMPUGEN INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      07-FEB-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      leukaemia),
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                                                                                                                                                                                                                                                                                                         present invention describes oligonucleotide libraries
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1225 CAAACCTCATTACCCACCAGAGGGCGCACACTGGGGAGAAGCCTTA 1270
                                                                                                                                                                                                                                                                                                                                                                                                                                           2002-257383/30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        variant;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        mouse;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity
                                                                                                                                                                                                                                                                                                                                         SEQ
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2001US-287724P
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                                                                                                                                                                                                                                                                                                                                    ID 6302; 47pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        transcriptome; oligonucleotide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DNA;
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71.78;
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 this patent did
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); Mismatches
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form part of
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                            mice,
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the printed
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                                                                                                                                                                                                                                                                                                       detecting
                                                                                                                                                                                                                                                                                                                                                                                             unit of
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RESULT 32
ABN45438
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
                                biological sample, in expression profiling studies, in qualitatively or quantitatively characterising the corresponding transcriptome, and in detecting RNA transcripts and splice variants of human or animal transcriptomes. The libraries may also be used as specialised mini libraries to detect transcripts of a sub-transcriptome under a particular biological or pathological state, and so allowing the detection of tissue and pathology-specific genes such as those genes only expressed in specific tissue under a specific pathological condition; to detect developmental specific genes; and to detect RNA transcripts and splice variants of a transcriptome of a patient suffering from a particular disorder. ABN27253 to ABN59589 represent oligonucleotide sequences from rats, humans and mice, which are used in the examplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New oligonucleotide libraries comprising oligonucleotides which selectively hybridize to mRNAs transcribed from a transcription unit of a genome, useful for detecting tissue-, pathology-, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          specification, but was obtained in electronic format directly from WIPO
at ftp.wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                                                                           set of messenger RNAs transcribed from a given transcription unit of
the genome, which encodes one or more messenger RNA splice variants.
The oligonuclectide libraries are useful for detecting mRNAs from a
                                                                                                                                                                                                                                                                                                                    transcription units that populate a genome. The library comprises several oligonucleotides, each capable of hybridising selectively
                                                                                                                                                                                                                                                                                                                                                     The present invention describes oligonucleotide libraries for detecting messenger RNAs that populate a (sub-)transcriptome, where the (sub-)transcriptome comprises messenger RNAs transcribed from multiple
                                                                                                                                                                                                                                                                                                                                                                                           The present invention describes oligonucleotide libraries
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           28-JUL-2000;
02-MAY-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             splice variant;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABN45438
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABN45438 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    developmental-specific genes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Shoshan A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 20-JUL-2001; 2001WO-IB01903
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO200210449-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human spliced transcript detection oligonucleotide SEQ ID NO:18186
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         15-JUL-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local
exemplification of the present invention.

The sequence data for this patent did not form part of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1225 CAAACCTCATTACCCACCAGAGGGCGCACACTGGGGAGAAGCCTTATGTTTGCA 1278
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2002-257383/30
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ن
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 l Similarity 66.7
36; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CACACCTGACTCAGCATCAGAGGGTTCATACTGGAGAGACCTTTCGAATGTA 58
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          60
                                                                                                                                                                                                                                                                                                                                                                                                                              SEQ ID 18186; 47pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Wasserman A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          B₽;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2000US-221607P
2001US-287724P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           rat; splice transcript; detection; RNA tr
t; transcriptome; oligonucleotide library;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Α,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          13 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                60
                                                                                                                                                                                                                                                                                                   each capable of hybridising selectively to a nscribed from a given transcription unit of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mintz E,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          14 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 25.2; D
Pred. No. 1.4e
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                               English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        16 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mintz L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  .4e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Faigler
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RNA transcript
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     60;
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0;
   the printed
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AAS18907 ID AAS1 XX

AAS18907

standard; DNA;

41

ВP

RESULT 34

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RESULT 33
ABL56733
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Best Local
                                                                 Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
                                                     Matches
                                                                                                                                                                                                                                                               Human Krupple Zinc Finger related protein 70.84 and encoding polynucleotide, useful in the diagnosis and treatment of malignant tumours, haemal disorders, human immunodeficiency virus infection,
                                                                                                       Sequence 41
                                                                                                                                             krupple zinc finger related protein 70.84. The polypeptide and it encoding polynucleotide are used in diagnosis and treatment of malignant tumour, haemal disorders, human immunodeficiency virus infection, immunological diseases and various inflammations. The
                                                                                                                                                                                                                                                                                                                                                                                                      07-JUL-2000; 2000CN-0117012
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO200226807-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       gene therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human; krupple zinc finger related haemal disorder; HIV infection; imm
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Probe #1 for krupple zinc finger related protein 70.84
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              specification, but was obtained in electronic format
at ftp.wipo.int/pub/published_pct_sequences.
                                                                                                                                 polynucleotide may also be used
                                                                                                                                                                                                Probes ABL56733-34 were used to identify cDNA encoding a human
                                                                                                                                                                                                                            Example 7; Page 15; 39pp; Chinese.
                                                                                                                                                                                                                                                                                                                          WPI;
                                                                                                                                                                                                                                                                                                                                                    мао Y,
                                                                                                                                                                                                                                                                                                                                                                            (BIOW-) BIOWINDOW GENE DEV INC SHANGHAI
                                                                                                                                                                                                                                                                                                                                                                                                                                 02-JUL-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                           04-APR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        30-JUL-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABL56733;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABL56733 standard; DNA; 41
                                                                                                                                                                                                                                                       immunological diseases
                                                                Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1192 GCAAGGAATGTGGGCAGAGCTTTAGCCTGAAGTCAAACCTCATTACCCCACCAGA 1245
                                                                                                                                                                                                                                                                                                                       2002-292482/33
             301 TGCTGGAGACTTATAACCATCTGGTCTCACTGGAAATTCCA 341
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 l Similarity
36; Conserv
                                                     31;
                                                                 Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GCAATGAACTGGGGAAAACCTTTCACTGTAAGTCATTCCCTCACTGTCCATCAGA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    60 BP; 19 A;
TGCTGGAGAATTATGGGAATGTGGCCTCCCTGGGATTTCCA 41
                                                                                                       BP; 8 A; 8 C; 13 G; 12 T; 0 other;
                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                 2001WO-CN01114
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        probe; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0.9%;
                                                                 75
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                                                                                                                                                                                                                                                       and
                                                                .
6%;
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                                                                                                                                                                                                                                                        inflammation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0;
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                                                   0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 25.2;
Pred. No. 1
                                                                Pred.
                                                                           Score
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                for gene therapy.
                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ied protein 70.84; tumo
immunological disease;
                                                              25; DB 24;
No. 1.3e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             4e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          24;
                                                                           Length 41;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   tumour
                                                   Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             directly from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CDNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                  Gaps
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RESULT 35
ABN33912
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                                                                                                                                                                                                                                                                                                                                                                                              Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to a new polypeptide-OZF (not defined) gene related protein 13, the polynucleotide encoding it, the process for preparing the said polypeptide by DNA recombination, the application of said polypeptide in treating several diseases such as cancer and HIV infect.on (human immunodeficiency virus). The antagonist against the said polypeptide and its therapeutic action, and the application of said polynucleotide for coding this new-OZF correlated protein 13 are disclosed. The present sequence is an oligonucleotide probe used to
28-JUL-2000; 2000US-221607P
                                                                                                                               splice variant;
                                                                                                                                                                 Human spliced transcript detection oligonucleotide SEQ ID NO:6660
                         20-JUL-2001; 2001WO-IB01903
                                                   07-FEB-2002
                                                                            WO200210449-A2
                                                                                                                                                                                                15-JUL-2002
                                                                                                                                                                                                                                                 ABN33912 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 41 BP; 14 A; 7 C; 11 G; 9 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Polypeptide-OZF gene related protein 13 and it -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2002-056303/08
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       27-MAR-2000; 2000CN-0115166
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (BODE-) BODE GENE DEV CO LTD SHANGHAI.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human OZF gene related protein 13 oligonucleotide probe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           26-MAR-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAS18907;
                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity
                                                                                                                                                                                                                                                                                                                             1411 TAAGGACACACACAGGAGAGAGCCTTATGTATGCACAGAA 1451
                                                                                                                                        mouse; rat; splice transcript; detection; RNA transcript
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ss; OZF gene related protein 13; cancer; HIV infection; immunodeficiency virus; probe.
                                                                                                                                                                                                                                                                                                                  TGAGAAGTCACACAGGTGAGCAGCCCTATGTATGTAATGAA 41
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Page 20; 31pp; Chinese.
                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                              acids encoding OZF gene related protein 13.
                                                                                                                            transcriptome; oligonucleotide library;
                                                                                                                                                                                                                                                 DNA;
                                                                                                                                                                                                                                                                                                                                                                               0.98;
                                                                                                                                                                                                                                                                                                                                                                    0,
                                                                                                                                                                                                                                                                                                                                                                   Score 25; DB
Pred. No. 1.3e
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                             DB 24;
                                                                                                                                                                                                                                                                                                                                                                                 3e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   polynucleotide encoding
                                                                                                                                                                                                                                                                                                                                                                   10;
                                                                                                                                                                                                                                                                                                                                                                                            Length 41;
                                                                                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                   0
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RESULT 36
ABN33916
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   δÃ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CC transcription units that populate a genome. The library comprises conserved the genome, which encodes one or more messenger RNAs transcribed from a given transcription unit of the genome, which encodes one or more messenger RNA splice variants.

CC The oligonucleotide libraries are useful for detecting mRNAs from a composition of the genome, which encodes one or more messenger RNA splice variants.

CC The oligonucleotide libraries are useful for detecting mRNAs from a composition of the printed libraries are useful for detecting mRNAs from a composition of the printed libraries and splice variants of human or animal composition.

CC transcriptomes. The libraries may also be used as specialised minical libraries to detect transcripts of a sub-transcriptome under a composition of tissue-and pathological state, and so allowing the condition; to detect developmental specific genes such as those genes conly expressed in specific tissue under a specific pathological condition; to detect developmental specific genes; and to detect RNA composition and splice variants of a transcriptome of a pathent suffering composition and pathological state, and so allowing the condition; to detect developmental specific genes; and to detect RNA composition and splice variants of a transcriptome of a pathent suffering composition of the present invention.

CC manual pathological state, and so allowing the condition of the present invention.

CC manual pathological state, and so allowing the specific pathological state, and so allowing the condition of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Best Loc
Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
                                 20-JUL-2001; 2001WO-IB01903
                                                                          07-FEB-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                at ftp.wipo.int/pub/published_pct_sequences.
                                                                                                                WO200210449-A2
                                                                                                                                                        Homo sapiens
                                                                                                                                                                                               splice
                                                                                                                                                                                                                    Human;
                                                                                                                                                                                                                                                       Human spliced transcript detection oligonucleotide SEQ
                                                                                                                                                                                                                                                                                                 15-JUL-2002
                                                                                                                                                                                                                                                                                                                                         ABN33916
                                                                                                                                                                                                                                                                                                                                                                            ABN33916 standard; DNA; 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New oligonucleotide libraries comprising oligonucleotides which selectively hybridize to mRNAs transcribed from a transcription a genome, useful for detecting tissue-, pathology-, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 1; SEQ ID 6660; 47pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               developmental-specific genes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2002-257383/30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 02-MAY-2001; 2001US-287724P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (sub-)transcriptome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               messenger RNAs that
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1033 GTGGGCGAGGCTTTACGTGGAAGTCAAACCTGATCACACATCAGAGGACACACTCAGGGG 1092
                                                                                                                                                                                                                mouse; rat; splice transcript; detection; RNA transcript;
                                                                                                                                                                                               variant;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               l Similarity
38; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GTGGCAAAAGCTTTGGAAGGACTTCCCATCTAAGCCAACATCAGCGTATTCACACAGGGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               invention describes oligonucleotide libraries
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Wasserman
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BP; 18 A; 15 C;
                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                             transcriptome;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.98;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    on describes oligonucleotide libraries for detecting populate a (sub-)transcriptome, where the comprises messenger RNAs transcribed from multiple
                                                                                                                                                                                                                                                                                                                                                                            ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mintz E,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          15 G; 12 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 24.8; E
Pred. No. 1.9e
0; Mismatches
                                                                                                                                                                                             oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mintz L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                .9e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Faigler
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length
                                                                                                                                                                                                                                                         ID NO:6664
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   60;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    unit of
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28-JUL-2000; 2000US-221607P

Of.

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                                                                                                                                                                                                                                                                                                                                  RESULT 37
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     quantitatively characterising the corresponding transcriptome, and in detecting RNA transcripts and splice variants of human or animal transcriptomes. The libraries may also be used as specialised mini libraries to detect transcripts of a sub-transcriptome under a particular biological or pathological state, and so allowing the detection of tissue- and pathology-specific genes such as those genes only expressed in specific tissue under a specific pathological condition; to detect developmental specific genes; and to detect RNA transcripts and splice variants of a transcriptome of a pattient suffering from a particular disorder. ABN27253 to ABN9589 represent oligonucleotide sequences from rats, humans and mice, which are used in the exemplification of the present invention.

N. B. The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New oligonucleotide libraries comprising oligonucleotides which selectively hybridize to mRNAs transcribed from a transcription a genome, useful for detecting tissue-, pathology-, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        transcription units that populate a genome. The library comprises several oligonucleotides, each capable of hybridising selectively to set of messenger RNAs transcribed from a given transcription unit of the genome, which encodes one or more messenger RNA splice variants. The oligonucleotide libraries are useful for detecting mRNAs from a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               biological sample, in expression profiling studies, in qualitatively or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       messenger RNAs that populate a (sub-)transcriptome, where the (sub-)transcriptome comprises messenger RNAs transcribed from multiple
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      developmental-specific genes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2002-257383/30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 02-MAY-2001; 2001US-287724P
                                                                                                                                                                                                                                                                                                  ABN34111 standard; DNA; 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 1;
28-JUL-2000;
                             20-JUL-2001; 2001WO-IB01903
                                                                                          WO200210449-A2
                                                                                                                                                        splice
                                                                                                                                                                                                      Human spliced
                                                                                                                                                                                                                                     15-JUL-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (COMP-) COMPUGEN INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ftp.wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                                                                                                                                                                                                                            1659 CACCAGAGGACACTCAGGGGAGAGCCATTTGTATGTGCTGAGTGTGGACGAGGCTTT
                                                                                                                                                      mouse; rat; splice transcript; detection; RNA tr
variant; transcriptome; oligonucleotide library;
                                                                                                                                                                                                                                                                                                                                                                                                                                               38;
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity
                                                                                                                                                                                                                                                                                                                                                                                CATCAGATAACCCACTTAGAAGAGAAACAATGTAAATGTGATGTATATGGCAAGGTATTT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              60 BP; 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQ ID 6664; 47pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          invention
                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
2000US-221607P
                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                    transcript detection oligonucleotide SEQ ID NO:6859
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Α;
                                                                                                                                                                                                                                                                                                                                                                                                                                                            0.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Ą
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              8 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        describes oligonucleotide libraries for detecting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mintz E,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            12 G;
                                                                                                                                                                                                                                                                                                  ВÞ
                                                                                                                                                                                                                                                                                                                                                                                                                                               0;
                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 24.8; I
Pred. No. 1.9e
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            16 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mintz L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                            1.9e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DΒ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Faigler
                                                                                                                                                                     RNA transcript;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length
                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              60;
                                                                                                                                                                                                                                                                                                                                                                                                                                             0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     unit
                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     of
f
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                                                                                                                                                                                                                                                                                                                                                                                                                1718
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망
                                                                                                                                                                                                                                                                                                                 ABA03751
                                                                                                                                                                                                                                                                                                  ID
                                                                                                                                                                                                                                                                                                                                RESULT 38
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CN1307039-A

Homo sapiens

26-JAN-2000; 08-AUG-2001.

2000CN-0111545

HIV;

n; zinc finger protein 22; cancer; nosohaemia; infection; human immunodeficiency virus; immunological disease;

inflammatory Human; zinc

disorder;

Human zinc finger protein

22 DNA probe

ABA03751;

18-FEB-2002

(first entry)

ABA03751 standard; DNA; 41

ВΡ

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0;

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Ωy
                                                                                                                                                                                                      the genome, which encodes one or more messenger RNA splice variants.

Control the oligonucleotide libraries are useful for detecting mRNAs from a biological sample, in expression profiling studies, in qualitatively or quantitatively characterising the corresponding transcriptome, and in the corresponding transcriptome. The libraries may also be used as specialised minical libraries to detect transcripts of a sub-transcriptome under a particular biological or pathological state, and so allowing the corresponding transcriptome under a gerticular biological or pathological state, and so allowing the condition; to detect developmental specific genes such as those genes conjy expressed in specific tissue under a specific pathological condition; to detect developmental specific genes; and to detect RNA cranscriptome of a particular disorder. ABN37253 to ABN59589 represent coligonucleotide sequences from rats, humans and mice, which are used in the exemplification of the present invention.

Construction of the present invention.

Construction but was obtained in electronic format directly from WIPO at fifth the problem of the printed specification, but was obtained in electronic format directly from WIPO at the problem of the printed control of the present invention.
                                                                                          Matches
                                                                                                                               Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    messenger RNAs that populate a (sub-)transcriptome, where the (sub-)transcriptome comprises messenger RNAs transcribed from multipl transcription units that populate a genome. The library comprises several oligonucleotides, each capable of hybridising selectively to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New oligonucleotide libraries comprising oligonucleotides which selectively hybridize to mRNAs transcribed from a transcription a genome, useful for detecting tissue-, pathology-, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 1; SEQ ID 6859; 47pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        02-MAY-2001; 2001US-287724P
                                                                                                                                                                        Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  set of messenger RNAs transcribed from a given transcription unit
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present invention describes oligonucleotide libraries
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          developmental-specific
                                                                                                           Locar
                         204 GAGGCGTTCGTGGCATTCAGGGATGTGGCTGTGTACTTCACCCAGGAGGAGT 255
                                                                                          ა
5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUGEN INC
                                                                                                           Similarity
                                                                                                                                                                          60
GAGGAATTTGTGACCCTCAAGGATGTCGGCATGGACTTCACCTTGGGAGACT
                                                                                        Conservative
                                                                                                                                                                          BP; 13
                                                                                                                                                                          A
                                                                                                     0.98;
                                                                                                                                                                        14 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          genes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mintz E,
                                                                                        0
                                                                                                                                                                        18 G; 15 T; 0 other;
                                                                                    Pred. No. 1.96
0; Mismatches
                                                                                                       Score 24.8;
Pred. No. 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mintz L,
                                                                                                           .9e+03;
                                                                                                                                   DB
                                                                                                                             24;
                                                                                        17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Faigler
                                                                                                                               Length 60;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        for
                                                                                        0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               multiple
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        detecting
                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   unit
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26-JAN-2000; 2000CN-0111545

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RESULT 39
ABA03752
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           멍
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Q
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
                                                                                      The invention relates to a novel polypeptide, human zinc finger protein 22. The polypeptide is useful for treating various diseases, such as malignant tumours, nosohaemia, HIV infection, immunological diseases and
     Sequence 41 BP;
                                                                                                                                                                                                                                                Polypeptide-human zinc finger protein 22 and said polypeptide -
                                        inflammatory disorder. The present sequence is a probe used to detect polynucleotides encoding the polypeptide of the invention
                                                                                                                                                                                                   Example 7;
                                                                                                                                                                                                                                                                                                                          WPI; 2002-062741/09
                                                                                                                                                                                                                                                                                                                                                                            Мао Ү,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     26-JAN-2000; 2000CN-0111545
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  26-JAN-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CN1307039-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human; zinc finger protein 22; cancer; nosohaemia; infection; HIV; human immunodeficiency virus; immunological disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABA03752 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        08-AUG-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   18-FEB-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABA03752;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention relates to a novel polypeptide, human zinc finger protein 22. The polypeptide is useful for treating various diseases, such as malignant tumours, nosohaemia, HIV infection, immunological diseases and inflammatory disorder. The present sequence is a probe used to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                inflammatory disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 41 BP; 17 A; 8 C; 11 G; 5 T; 0 other; .
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 7; Page 20 (disclosure); 34pp; Chinese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Polypeptide-human zinc finger protein 22 and said polypeptide - \,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         detect polynucleotides encoding the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (BODA-) BODAO GENE TECH CO LTD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1066 TCACACATCAGAGGACACTCAGGGGAGAAACCTTATG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2002-062741/09
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            zinc finger protein 22 DNA probe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        _
                                                                                                                                                                                                                                                                                                                                                                          Xie
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      30;
                                                                                                                                                                                                                                                                                                                                                                                                                  BODAO GENE TECH CO LTD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Xie
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Page 20 (disclosure); 34pp; Chinese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2000CN-0111545
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
  16 A; 9 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.9%;
76.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   probe; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 41
  11 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 24.6;
Pred. No. 1
                                                                                                                                                                                                                                                                                                                                                                                                                     SHANGHAI.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SHANGHAI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
  5 T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         polypeptide of
0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1.7e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            . 24;
                                                                                                                                                                                                                                                                    polynucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         polynucleotide for coding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              9;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1104
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              41;
                                                                                           diseases and
                                                                                                                                                                                                                                                                    for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0
                                                                                                                                                                                                                                                             coding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0,;
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RESULT 41
AAH89253
ID AAH89
XX
AC AAH89

AAH89253 standard; DNA; 51

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAH89252
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 40
                                                                                                Matches
                                                                                                                     Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Best Local
                                                                                                                                                                                                                   abberrant protein expression, including cancer, amyloidosis, diabetes, Alzheimer's disease, Down's syndrome, oedema, lupus (SLE), vasculitis, glomerulonephritis, haemolytic anaemia, thrombocytopaenia, arthritis, meningitis, muscular disorders, dementia, neurological diseases, tuberc sclerosis, male infertility, hypercalcaemia, blood pressure disorders, osteoporosis, pathogenic infections, hypercholesterolaemia, obesity or autoimmunity. The present sequence is a polymorphism-containing oligonucleotide fragment of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                         The present invention relates to human nucleic acids containing single nucleotide polymorphisms (SNPs). These can be used in forensic and paternity tests, and relating the treatment of diseases associated with the containing the co
                                                                                                                                                                                        Sequence 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Shimkets RA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   07-JAN-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 05-JAN-2001; 2001WO-US00322
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          P-PSDB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               19-JUL-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO200151670-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; single nucleotide polymorphism; SNP; forensic test; aberrant protein expression:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human transcription
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      01-OCT-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAH89252;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAH89252 standard;
                   1251 CACACTGGGGAGAAGCCTTATGTTTGCAGGGAATGTGGGCGTGGCTT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1066 TCACACATCAGAGGACACACTCAGGGGAGAAACCTTATG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2001-451871/48
                                                                                                                   Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         30;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAM00143.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity
CACACTGGAGAGAACCCTATAGATGTTGTGGATGTGGGAAGGCCTT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Page 119; 475pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TGAGACATCAGAGAGTGCACCCAGGAGAGAAACCTTACG
                                                                                                                                                                                      BP; 13 A;
                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2000US-0174962
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Leach
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                aberrant protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DNA;
                                                                                                                0.98;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ¥
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      factor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0.98;
                                                                                                                                                                                      10 C; 16 G; 12 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DNA polymorphic site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ВР
                                                                                             0
                                                                                                            Score 24.6; DB Pred. No. 2e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 24.6;
Pred. No. 1
                                                                                        Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              expression;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            .7e+03
                                                                                                                                                                                                                                                                                                                        neurological diseases, tuberous
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DΒ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                paternity ds.
                                                                                           14;
                                                                                                                                    22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    9;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           39
                                                                                        Indels
                                                                                                                                 Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NO:
50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 41;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               cancer,
                                                                                      Gaps
                                                                                        0
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RESULT 42
AAD33628/c
                                                                                                                                                                                                                                                                      B
                                                                                                                                                                                                                                                                                            QУ
                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local
                                                                                                                                                                                                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                                                                                                                                           aberrant protein expression, including cancer, amyloidosis, diabetes, Alzheimer's disease, Down's syndrome, oedema, lupus (SLE), vasculitis, glomerulonephritis, haemolytic anaemia, thrombocytopaenia, arthritis, meningitis, muscular disorders, dementia, neurological diseases, tuberous sclerosis, male infertility, hypercalcaemia, blood pressure disorders, osteoporosis, pathogenic infections, hypercholesterolaemia, obesity or autoimmunity. The present sequence is a polymorphism-containing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present invention relates to human nucleic acids containing single nucleotide polymorphisms (SNPs). These can be used in forensic and paternity tests, and to aid in the treatment of diseases associated with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Shimkets RA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        07-JAN-2000; 2000US-0174962.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 05-JAN-2001; 2001WO-US00322
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human; single nucleotide polymorphism; SNP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     01-OCT-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 1; Page 119; 475pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      forensic
                                                                                                                                                    01-JUL-2002
                                                                                                                                                                                                                                                                                                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                                                                                  oligonucleotide fragment of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Isolated human polynucleotides containing single nucleotide polymorphisms, useful for the treatment and diagnosis of e.g. cancer,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   P-PSDB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (CURA-) CÚRAGEN CORP
            31-JAN-2002
                                                             Unidentified
                                                                                                 DNA binding protein;
                                                                                                                           Zif-3
                                                                                                                                                                                                       AAD33628
                                    WO200208286-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                infection and
                                                                                                                                                                                                                                                                                                                      Local Similarity
mes 33; Conserv
                                                                                                                                                                                                                                                                                           1251 CACACTGGGGAGAAGCCTTATGTTTGCAGGGAATGTGGGCGTGGCTT 1297
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2001-451871/48
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            transcription
                                                                                                                         specific antisense oligo used to generate three finger ZFP
                                                                                     replication;
                                                                                                                                                                                                                                                                      4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAM00144.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      test;
                                                                                                                                                                                                                                                                                                                                                                          51 BP; 14 A;
                                                                                                                                                                                                       standard;
                                                                                                                                                                                                                                                                      CACACTGGAGAGAAACCCTATAAATGTTGTGGATGTGGGAAGGCCTT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Leach MD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                diabetes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      aberrant protein
                                                                                     gene expression; virucide; Zif3;
                                                                                                                                                                                                       DNA;
                                                                                                                                                                                                                                                                                                                                 0.9%;
                                                                                                 zinc finger domain 3;
                                                                                                                                                                                                                                                                                                                                                                       10 C; 15 G; 12 T; 0 other;
                                                                                                                                                                                                       55
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DNA
                                                                                                                                                                                                       ΒP
                                                                                                                                                                                                                                                                                                                       0;
                                                                                                                                                                                                                                                                                                                     Score 24.6; D
Pred. No. 2e+0
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                  the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          polymorphic site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     expression;
                                                                                                                                                                                                                                                                                                                                  e 24.6; DB 2
. No. 2e+03;
                                                                                                                                                                                                                                                                                                                                               DB 22;
                                                                                                 zinc finger protein; ZFP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        paternity test; ds.
                                                                                                                                                                                                                                                                                                                       14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQ
                                                                                                                                                                                                                                                                                                                       Indels
                                                                                                                                                                                                                                                                                                                                               Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ID NO:
                                                                                                                                                                                                                                                                      50
                                                                                                                                                                                                                                                                                                                                                  51;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             34
                                                                                                                                                                                                                                                                                                                      0;
                                                                                                                                                                                                                                                                                                                       Gaps
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RESULT 43
ABN34011
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Qy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              virus), modulating gene expression, detecting an altered zinc finger recognition sequence and diagnosing disease associated with abnormal genomic structure. The present DNA sequence is zinc finger domain 3 (Zif-3) specific oligonucleotide which is used to generate three finger ZFP for the L1 site of beet curly top virus (BCTV). This sequence is used in the exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention relates to novel DNA binding proteins comprising zinc finger domains in which two histidine and two cysteine residues coordinate a central zinc ion. The invention particularly relates to
                                                                                                                                                                                                                                                                                                                                                        ABN34011;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      the identification of a context-independent recognition code to design zinc finger domains. The invention also relates to zinc finger proteins (ZFP) designed using this recognition code. The ZFPs are useful for altering genomic structure, inhibiting viral replication (where viral
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New Zinc Finger Protein (ZFP) comprising three essential domains us for diagnosing diseases associated with abnormal genomic structure
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (SYGN)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      19-JUL-2001;
                                                                                                                 28-JUL-2000;
02-MAY-2001;
                                                                                                                                                                                 07-FEB-2002
                                                                                                                                                                                                           WO200210449-A2
                                                                                                                                                                                                                                    Homo sapiens
                                                                                                                                                                                                                                                            splice
                                                                                                                                                                                                                                                                                                   Human spliced
                                                                                                                                                                                                                                                                                                                                                                                  ABN34011 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        for diagnosing
                                                                                                                                                       20-JUL-2001;
                                                                                                                                                                                                                                                                                                                               15-JUL-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            replication is inhibited for plant virus, an animal virus or a human
                                                                                        (COMP-) COMPUGEN INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1211 CTTTAGCCTGAAGTCAAACCTCATTACCCACCAGAGGGGGGGCACACTGGGGAGAAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2002-172000/22
                                      2002-257383/30
                                                                                                                                                                                                                                                                                                                                                                                                                                                  55
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     l Similarity
36; Conserv
                                                                                                                                                                                                                                                                          mouse; rat; splice transcript;
                                                                                                                                                                                                                                                               variant;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SYNGENTA PARTICIPATIONS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              7; Page 72;
                                                              Α,
                                                                                                                                                                                                                                                                                                                                                                                                                                                 55 BP; 9 A; 14 C; 15 G; 17 T; 0 other;
                                                                Wasserman
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2000US-220060P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2001WO-EP08367
                                                                                                                2000US-221607P
2001US-287724P
                                                                                                                                                         2001WO-IB01903
                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                    transcript detection oligonucleotide
                                                                                                                                                                                                                                                              transcriptome;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              143pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (ZFP) comprising three essential domains useful
                                                                                                                                                                                                                                                                                                                                                                                   60
                                                                Mintz
                                                                                                                                                                                                                                                                                                                                                                                   ΒP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 24.6; D
Pred. No. 2.1e
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0;
                                                                                                                                                                                                                                                            oligonucleotide library;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ΑG
                                                                E
                                                                Mintz
                                                                                                                                                                                                                                                                          detection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  .le+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 24;
                                                                Ľ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      19;
                                                                Faigler
                                                                                                                                                                                                                                                                           RNA transcript;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length
                                                                                                                                                                                                                                                                                                     SEQ
                                                                                                                                                                                                                                                                                                     ID NO:6759
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                55;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1265
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New oligonucleotide libraries comprising oligonucleotides which selectively hybridize to mRNAs transcribed from a transcription  ${\bf r}$ 

unit

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A A C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C A C C 
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       the genome, which encodes one or more messenger RNA splice variants.

CC The oligonucleotide libraries are useful for detecting mRNAs from a guantitatively characterising the corresponding transcriptome, and in CC quantitatively characterising the corresponding transcriptome, and in CC transcriptomes. The libraries may also be used as specialised mini CC libraries to detect transcripts of a sub-transcriptome under a CC quarticular biological or pathological state, and so allowing the CC only expressed in specific tissue under a specific pathology as those genes condition; to detect developmental specific genes such as those genes condition; to detect developmental specific genes; and to detect RNA CC from a particular disorder. ABN27253 to ABN39589 represent the exemplification of the present invention.

CC N.B. The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPD)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 44
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local
                                                                    28-DEC-1999; 99US-0173419.
27-DEC-2000; 2000US-0173419.
                                                                                                                                                                                                                                                                                                                                                                                                                                   Immunosuppressive; immunostimulatory; antiinflammatory; cytostatic; neuroprotective; antimicrobial; gene therapy; vaccine; amylase; canc amyloid protein; angiopoietin; apoptosis related protein; cadherin; cyclin; polymerase; oncogene; histone; kinase; colony stimulating fa complement related protein; cytochrome; kinesin; cytokine; interfero interleukin; G-protein coupled receptor; thioesterase; inflammation;
                    (CURA-) CURAGEN CORP
                                                                                                                                                                                                                                                                                    WO200147944-A2
                                                                                                                                                                   28-DEC-2000;
                                                                                                                                                                                                                            05-JUL-2001.
                                                                                                                                                                                                                                                                                                                                          Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                multifactorial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human SNP oligonucleotide #5867
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  24-JAN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAL32659;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAL32659 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 transcription units that populate a genome. several oligonucleotides, each capable of he set of messenger RNAs transcribed from a cit
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention describes oligonucleotide libraries for detecting messenger RNAs that populate a (sub-)transcriptome, where the (sub-)transcriptome comprises messenger RNAs transcribed from multiple
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     a genome, useful for de developmental-specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 1; SEQ ID 6759; 47pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ftp.wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            eral oligonucleotides, each capable of hybridising selectively to of messenger RNAs transcribed from a given transcription unit of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              G
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAGTCTTGAGCCATCATTCATCCCTTAACAGGCACATTAGATCTTACACCAGACA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BP;
                                                                                                                                                             2000WO-US35498
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                            disease; ss
                                                                                                                                                                                                                                                                                                                                                                                                                          disease; autoimmune
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         20 A; 17 C; 9 G; 14 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       detecting tissue-,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     genes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Pred. No. 2.200; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 24.6;
Pred. No. 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           :.2e+03;
nes 19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       pathology-,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The library comprises
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                60;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    interferon;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     for detecting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       cancer,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             59
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           factor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0;
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AAL33200
ID AAL3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 45
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CC The present invention relates to oligonucleotides encoding polymorphic cc variants of proteins related to amylases, amyloid proteins, angiopoietin, appropriate of proteins, cadherin, cyclin, polymerase, oncogenes, cc apoptosis related proteins, cadherin, cyclin, polymerase, oncogenes, cc proteins, kinases, colony stimulating factors, complement related cc proteins, cytochromes, kinesins, cytokines, interferons, interleukins, cc one such oligonucleotide. The oligonucleotides and the pertiades encoded cc by them may be used in the prevention, diagnosis and treatment of cd iseases associated with inappropriate expression of the proteins listed cc above. Disorders that may be prevented, diagnosed and/or treated include cc diseases (e.g. rheumatoid arthritis, multiple sclerosis, diabetes, cs ystemic lupus erythromaticsus and Grave's diseases, inflammation, cancer cc (e.g. cancers of the bladder, brain, breast, colon and kidney, leaked cc ordanisms), diseases of the nervous system and an infection of pathogenic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local :
                          Shimkets RA,
                                                                                 28-DEC-1999;
27-DEC-2000;
                                                                                                                                                        05-JUL-2001
                                                                                                                                                                                                                                                                                      Immunosuppressive; immunostimulatory; antlinflammatory; cytostatic neuroprotective; antimicrobial; gene therapy; vaccine; amylase; ca amyloid protein; angiopoietin; apoptosis related protein; cadherin cyclin; polymerase; oncogene; histone; kinase; colony stimulating
                                                                                                                         28-DEC-2000; 2000WO-US35498
                                                                                                                                                                                    WO200147944-A2
                                                                                                                                                                                                                  Homo
                                                                                                                                                                                                                                                         multifactorial
                                                                                                                                                                                                                                                    comprement related protein; cytochrome; kinesin; cytokine; interferon; interleukin; G-protein coupled receptor; thioesterase; inflammation; multifactorial disease; autoimmune disease; infection;
                                                                                                                                                                                                                                                                                                                                                                    Human SNP oligonucleotide #6408
                                                                                                                                                                                                                                                                                                                                                                                                   24-JAN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                AAL33200
                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAL33200 standard; DNA; 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 1; Page 3073; 4143pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2001-465210/50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Shimkets RA,
                                                                                                                                                                                                               sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1353 TACATTTGCAGGGAGTGTGAGCAAGGCTTTAGCCAGAAGTCACACCTCAT 1402
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURAGEN
                                                                                                                                                                                                                                          system disease; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity
                                                                                                                                                                                                                                                                                related protein;
                                                                                99US-0173419
2000US-0173419
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ВP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                         Leach M;
                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                     CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Leach
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Α;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            σ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        ΒP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         14 G; 16 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 24.4; DB 22
Pred. No. 2.3e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
                                                                                                                                                                                                                                                                                                                                 antiinflammatory; cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length
                                                                                                                                                                                                                                                                                                          cadherin;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0,
                                                                                                                                                                                                                                                                                                                       cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                         factor;
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WPI; 2001-465210/50

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RESULT 46
ABN33573
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      by them may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate expression of the proteins listed above. Disorders that may be prevented, diagnosed and/or treated include multifactorial diseases with a genetic component, such as autoimmune diseases (e.g. rheumatoid arthritis, multiple sclerosis, diabetes, systemic lupus erythromatosus and Grave's disease), inflammation, cancer (e.g. cancers of the bladder, brain, breast, colon and kidney, leukaemia), diseases of the nervous system and an infection of pathogenic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention relates to oligonucleotides encoding polymorphic variants of proteins related to amylases, amyloid proteins, angiopoletin, apoptosis related proteins, cadherin, cyclin, polymerase, oncogenes, histones, kinases, colony stimulating factors, complement related
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  proteins, cytochromes, kinesins, cytokines, interferons, interleukins, G-protein coupled receptors and thioesterases. The present sequence is one such oligonucleotide. The oligonucleotides and the peptides encode
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Polymorphic nucleic acids encoding e.g. amylases, cyclins, polymerases, oncogenes and histones, useful for diagnosing and treating, e.g. cancer, autoimmune diseases and infections -
                                                                         New oligonucleotide libraries comprising oligonucleotides which selectively hybridize to mRNAs transcribed from a transcription a genome, useful for detecting tissue-, pathology-, and
                                                                                                                                                                                                                                      28-JUL-2000;
02-MAY-2001;
                                                                                                                                                                                                                                                                                                                  07-FEB-2002
                                                                                                                                                                                                                                                                                                                                                 WO200210449-A2
                                                                                                                                                                                                                                                                                                                                                                                                                            Human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human spliced transcript detection oligonucleotide SEQ ID NO:6321.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            15-JUL-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABN33573
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABN33573 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 51
The present invention describes oligonucleotide libraries
                                Example 1;
                                                                                                                                           WPI; 2002-257383/30
                                                                                                                                                                          Shoshan
                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                             splice variant;
                                                              developmental-specific
                                                                                                                                                                                                       (COMP-) COMPUGEN INC
                                                                                                                                                                                                                                                                                  20-JUL-2001; 2001WO-IB01903
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1548
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TTTAGCCTGAAGTCAAACCTTAACAAACACCAGAGGTCACACACGGGGGA 1597
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Page 3217; 4143pp; English
                             SEQ ID 6321; 47pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BP;
                                                                                                                                                                        Wasserman
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                     2000US-221607P
2001US-287724P
                                                                                                                                                                                                                                                                                                                                                                                                                            rat; splice transcript; detection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              16 A; 17 C; 10 G; 8 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                             transcriptome; oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0.98;
                                                                                                                                                                        A,
                                                              genes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        60
                                                                                                                                                                      Mintz E, Mintz L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ВP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 24.4;
Pred. No. 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 3e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  16;
                                                                                                                                                                      Faigler S;
                                                                                                                                                                                                                                                                                                                                                                                                             library;
                                                                                                                                                                                                                                                                                                                                                                                                                          RNA transcript
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length
for detecting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                           unit
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     encoded
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RESULT 47
ABN34042
DE KXXX PRI DE XXXX PRI DE XXX PRI DE XXXX PRI DE XXX PRI DE XXXX 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Вþ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                only expressed in specific tissue under a specific pathological condition; to detect developmental specific genes; and to detect RNA transcripts and splice variants of a transcriptome of a patient suffering from a particular disorder. ABN27253 to ABNS9589 represent oligonucleotide sequences from rats, humans and mice, which are used in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     quantitatively characterising the corresponding transcriptome, and in detecting RNA transcripts and splice variants of human or animal transcriptomes. The libraries may also be used as specialised mini
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            the genome, which encodes one or more messenger RNA splice variants. The oligonucleotide libraries are useful for detecting mRNAs from a biological sample, in expression profiling studies, in qualitatively
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   messenger RNAs that populate a (sub-)transcriptome, where the (sub-)transcriptome comprises messenger RNAs transcribed from multiple transcription units that populate a genome. The library comprises several oligonucleotides, each capable of hybridising selectively to a
                                                                                                                                             New oligonucleotide libraries comprising oligonucleotides which selectively hybridize to mRNAs transcribed from a transcription a genome, useful for detecting tissue-, pathology-, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                               28-JUL-2000;
02-MAY-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       07-FEB-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human; mouse; rat; splice transcript; detection; splice variant; transcriptome; oligonucleotide 1:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      the exemplification of the present invention. N.B. The sequence data for this patent did not form specification, but was obtained in electronic format
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     libraries to detect transcripts of a sub-transcriptome under a particular biological or pathological state, and so allowing the detection of tissue- and pathology-specific genes such as those genes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                set of messenger RNAs transcribed from a given transcription unit
                                                             Example 1;
                                                                                                                       developmental-specific
                                                                                                                                                                                                                                                                              WPI; 2002-257383/30
                                                                                                                                                                                                                                                                                                                                        Shoshan A, Wasserman A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         20-JUL-2001; 2001WO-IB01903
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO200210449-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human spliced transcript detection oligonucleotide SEQ ID NO:6790
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABN34042;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABN34042 standard; DNA; 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  15-JUL-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ftp.wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1746 CAGAGGACACATTCAGGGGAAAAAGCCTTTTATGTGCAGGGAGTGTGGCAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    11 CAGAGTCTTCACACAGTGGGAGTGTCATTTAAGTGCAGTGACTGTGGAAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           34;
                                                                                                                                                                                                                                                                                                                                                                                                        COMPUGEN INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity
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                                                      SEQ ID 6790; 47pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                               2000US-221607P
2001US-287724P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  11 C;
                                                                                                                       genes
                                                                                                                                                                                                                                                                                                                                        Mintz E,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  20 G; 14 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 24.4; DB 24; Pred. No. 2.5e+03;
                                                      English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              oligonucleotide library;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                        Mintz L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           16;
                                                                                                                                                                                                                                                                                                                                        Faigler
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             format
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RNA transcript;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length
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                                                                                                                                                                                                                                                                                                                                        s
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0;
                                                                                                                                                                                 unit of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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The present invention describes oligonucleotide libraries for detecting

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RESULT 48
AAH77419
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Best Local S
Matches 37
                                                                      Human zinc finger protein 32 and encoded polynucleotide, diagnosis and treatment of e.g. nervous system diseases, timor, hemopathy, human immunodeficiency virus infection,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           detecting RNA transcripts and splice variants of human or animal transcriptomes. The libraries may also be used as specialised mini libraries to detect transcripts of a sub-transcriptome under a particular biological or pathological state, and so allowing the detection of tissue- and pathology-specific genes such as those genes only expressed in specific tissue under a specific pathological condition; to detect developmental specific genes; and to detect RNA transcripts and splice variants of a transcriptome of a patient suffer from a particular disorder. ABN27253 to ABN59589 represent oligonucleotide sequences from rats, humans and mice, which are used it is communicated to the process of transcriptome.
  Example
                                                                                                                                                                                                                                                                                                                                              07-MAR-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; zinc finger protein 32; cancer; haemopathy; development disorder; HIV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human zinc finger protein 32 coding sequence probe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           05-NOV-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             several oligonucleotides, each capable of hybridising selectively to set of messenger RNAs transcribed from a given transcription unit of the genome, which encodes one or more messenger RNA splice variants. The oligonucleotide libraries are useful for detecting mRNAs from a biological sample, in expression profiling studies, in qualitatively quantitatively characterising the corresponding transcriptome, and in
                                                                                                                                                                                                                                                                                                                                                                                                 26-FEB-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                     13-SEP-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO200166731-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAH77419 standard;
                                                                                                                                                                                                                                                                                          (BIOW-) BIOWINDOW GENE DEV INC SHANGHAI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               the exemplification of the present invention.

N.B. The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  transcription units that populate a genome. several oligonucleotides, each capable of h
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   messenger RNAs that populate a (sub-)transcriptome, where the (sub-)transcriptome comprises messenger RNAs transcribed from multranscription units that populate a genome. The library comprises
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ftp.wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1337 TTCAGGAGAGAGCCTTACATTTGCAGGAGTGTGAGCAAGGCTTTAGCCAGAAGTCA 1394
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              l Similarity
37; Conser
  7;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TACTGGAGAGCAACCCTACATATGTGAAGAATGTGTGAAATTTTTTTACCCAATCCTCA 58
Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   вP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                              2000CN-0111915
                                                                                                                                                                                                                                                                                                                                                                                               2001WO-CN00289
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  15;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A; 13 C;
34pp; Chinese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.98;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             41
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           probe;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0,;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                10 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              17 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SS
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No. 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            nervous system disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    .5e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      immune
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Indels
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                                                                         malignant
, and ·
                                                                                                                             used in
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The

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invention

provides the

protein

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coding

sequences of

human

Sequence 51 BP;

15 A;

13 C;

17 G; 6 T; 0 other,

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                         cc polynucleotide sequences. The sequences can be used in gene and protein claims by the control of therapy, and in vaccine production. (I) and the polypeptides encoded by them may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate expression of polymorphic polypeptides. For example, (I) may be used to treat disorders by rectifying mutations corrections in a patient's genome that affect the activity of polypeptides by expressing inactive proteins or to supplement the patients own production of polypeptide. Additionally, (I) and its complementary sequences may also be used as DNA probes in diagnostic assays to detect and quantitate the presence of similar nucleic acids in samples, and therefore which patients may be in need of restorative therapy. The polypeptides encoded by (I) may be used as antigens in the production of antibodies specific for polymorphic polypeptides. The antibodies may also be used to down regulate expression and activity. The antibodies may also be used as diagnostic agents for detecting the presence of polymorphic polypeptides in samples.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAI79772/c
ID AAI79772 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                                                                       sequences (I), which contain single nucleotide polymorphisms (SNPs). AAM53114 to AAM53329 represent peptides related to human polymorphic
                                                                                                                                                                                                                                                                                                                                                                             Claim 1; Page 2559; 2653pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2001-356160/37
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Shimkets
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          30-NOV-1999; 99US-0168138
29-NOV-2000; 2000US-0726173
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; single nucleotide polymorphism; SNP; gen protein therapy; vaccine; probe; diagnostic ass quantitation; restorative therapy; polymorphic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         zinc finger protein 32. The sequences can be used in the treatment of cancer, nervous system diseases, haemopathy, development disorders, HIV infection, immune diseases and inflammation. The present sequence is a probe for the coding sequence of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                 Polymorphic nucleic acid sequences, useful in genetic testing
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                                                                                                                                              Matches
                                                                                                                                                                                                                                               the identification of a context-independent recognition code to design zinc finger domains. The invention also relates to zinc finger proteins (ZFP) designed using this recognition code. The ZFPs are useful for altering genomic structure, inhibiting viral replication (where viral replication is inhibited for plant virus, an animal virus or a human virus), modulating gene expression, detecting an altered zinc finger recognition sequence and diagnosing disease associated with abnormal genomic structure. The present DNA sequence is zinc finger domain 3 (Zif-3) specific PCR primer 6 used to generate three finger ZFP. This sequence is used in the exemplification of the invention.
                                                                                                                                                                                                                  Sequence 55 BP; 7 A; 11 C; 11 G; 14 T; 12 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present invention relates to novel DNA binding proteins comprising zinc finger domains in which two histidine and two cysteine residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  21-Jul-2000; 2000US-220060P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DNA binding protein; zinc finger domain 3; zinc finger protein; ZFP; viral replication; gene expression; virucide; Zif3; PCR; primer; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 6; Page 71; 143pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New Zinc Finger Protein (ZFP) comprising three essential domains useful for diagnosing diseases associated with abnormal genomic structure -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (SYGN ) SYNGENTA PARTICIPATIONS AG
                                                                                                                                                                                                                                                                                                                                                                                                                                   coordinate a central zinc ion. The invention particularly relates to
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| 278<br>279<br>280<br>281<br>281<br>282<br>283   | 274<br>c 275<br>c 276<br>c 277  | c 271<br>c 272<br>273  | 268<br>c 269<br>270  | . 266<br>267                                 | 264   | 262<br>263                                       | 260<br>260   | 25 <b>8</b>  | 255<br>c 256  |  | c 251<br>c 252                                 | c 248<br>c 249<br>250   |  | 0 244                          |  |   | c 237   |  |  |   | c 230  | c 228                                  |                              | c 224                  |   |   | c 219  |  |                        | $\frac{213}{214}$      |
| 18.2<br>18.2<br>18.2<br>18.2<br>18.2  | 18.<br>18.  | 18.  | 18.<br>18.   | 18.<br>18.                                   | 18.   | 18.  | 18.  | 18.  | 18.<br>18.  | 18.<br>18.   | 18.  | 18.   | 18.  | 18.                            | 18.  |   |   | 18.  |  |   | 1100   | 18.                                    | 18.                          | <b>.</b>               | 18.<br>18.                              | 18.   | 18.  | ـــ د د  | 18.                    | 18.                    |
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|   |   |  |  |  |   |  |  |  |   |  |  |   |  |                                |  |   |   |  |  |   |  |  |                              |                        |   |   |  |  |                        |                        |
| AX158138<br>AX158434<br>AX159651<br>AX160010<br>AX160039<br>AX160041  | 142228<br>RHCHAN<br>AX010669<br>AX157231  | AX160040<br>AX160042<br>AX199546                                     | AX426599<br>E02126<br>AR148167                                       | AX322954<br>AX363268                         | AX282194<br>AX282303  | AR1/4681<br>AR165629                             | AX403352<br>A38852                                   | AX040753<br>AX195145                               | AX327057<br>AR127961                                    | A90731<br>AX456312                                   | AR091831<br>AR157739                           | CGPROMF75<br>AX369291   | AR054380<br>MUSIGHXR                               | A38984<br>AB054379             | AX164840<br>A69641   | AX161100<br>AX161690  | AX116225<br>AX157831  | 160387<br>AR086105                                       | AX162646   | A78547  | AX222266<br>HSA235211                                    | AR164192                               | AR144905                     | AX229635               | AR151117<br>AX424002                    | E09318  |  |  | AX189822               | AX165657               |
|   |   |  |  |  |   |  |  |  |   |  |  | Ţ.  | ~  |                                |  |   |   |  |  |   |  | •                                      |                              |                        |   |   | ā  | o  |                        |                        |
| AX<br>AX<br>AX  | I4<br>V<br>AX   | AX<br>AX   | AX<br>EO   | AX<br>AX                                     | AX<br>AX  | AR<br>AR   | AX<br>A3   | AX<br>AX   | AX<br>AR  | A9<br>AX   | AR<br>AR                                       | AX<br>AX  | AR<br>M  | A3<br>A3                       | A S  | AX<br>AX  | AX  | I6<br>AR   | AX:  | A71   | AJ.  | AR:                                    | AR.                          | AX                     | AR:                                     | EO  | S5.  | A  | , A                    | AX.                    |
| AX158138 Sequence<br>AX158434 Sequence<br>AX159651 Sequence<br>AX160010 Sequence<br>AX160039 Sequence<br>AX160041 Sequence  | 2228 Se<br>01206 V<br>010669<br>157231  | 160040<br>160042<br>199546   | 426599<br>2126 Pc<br>148167  | 322954<br>363268                             | 221438<br>282194<br>282303  | 174681<br>165629                                 | 403352<br>8852 Se                                    | 040753<br>195145                                   | 327057<br>127961  | 0731 Se<br>456312                                    | 091831   | 0365 C.<br>369291<br>002595   | 054380<br>13400 N                                  | 8984 SE                        | 164840<br>9641 Se  | 161100  | 116225<br>157831  | 0387 Se<br>086105  | 162646   | 1449 Se<br>8547 Se  | 235211   | 164192                                 | 144905                       | 229635                 | 151117<br>424002                        | 080003<br>9318 Pr   | 9925 TA  | F328257  | 189822                 | 165664                 |
| seque<br>seque<br>seque<br>seque  | eguenc<br>esicu<br>seque<br>seque   | seque<br>seque<br>seque  | Seque<br>)lylin<br>Seque   | Seque  | Seque   | Seque  | Seque  | Seque  | seque   | guenc  | Seque  | gluta<br>Seque  | Seque  | secue<br>Squenc                | Seque  | Seque   | Seque<br>Seque  | iquenc<br>Seque  | Seque  | quenc   | Seque  | Seque                                  | Seque                        | Seque                  | Seque                                   | Seque<br>imer.  | L1d1=  | Mus  | Seque                  | Seque                  |
| ince<br>ince  | llar s<br>nce<br>nce  | nce  | ker  | nce  | in ce   | nce  | 6 e  | nce  | nce   | e 15<br>nce  | ince<br>ince                                   | micu<br>nce   | ince<br>Ig ac                                      | e 71                           | 20<br>10<br>10<br>10<br>10<br>10<br>10<br>10<br>10<br>10<br>10<br>10<br>10<br>10 | nce   | nce   | nce  | nce  | 4 4   | sapi   | nce                                    | nce                          | nce                    | nce                                     | nce<br>9/1  | prot   | muscu  | nce                    |                        |
|   |   |  |  |  |   |  |  |  |   |  |  |   |  |                                |  |   |   |  |  |   |  |  |                              |                        |   |   |  |  |                        |                        |
|   |   |  |  |  |   |  |  |  |   |  |  |   |  |                                |  |   |   |  |  |   |  |  |                              |                        |   |   |  |  |                        |                        |
|   |   | <u>.</u> .   |  |  |   |  |  |  |   |  |  |   |  |                                |  |   |   |  | -  |   |  |  |                              | -                      |   |   |  |  |                        |                        |
|   | C 347<br>C 348<br>C 349   | 344<br>345   | 341<br>342<br>343  | 339  | . 336<br>337<br>338   | C 334  | C 332  | . 330<br>. 331                                     | 328<br>329  |  |  | c 321   |  |                                |  | c 312<br>313  |   | c 308.   | . c 306  | 305   | c 303  | C 300                                  | c 299                        | 297                    | C 295                                   | c 293   | 292  | c 290  | 288                    | 285                    |
| 351<br>352<br>352<br>354<br>355<br>1  | 347<br>348<br>349<br>350  | ا با با با   | بر بر بر   | ر بر بر                                      | ے نے نے   | 3354   | 332  | در در  | יין ניין  | 326 1<br>327 1                                       | 324 1<br>325 1                                 | 321 1<br>322 1  | 319 1<br>320 1                                     | 317<br>317<br>1                | 314  | 312 18.<br>313 18.  | 310 18.<br>311 18.  | 308. 18.<br>309 18.                                      | 306 18.<br>307 18.   | 305 18.   | 302 18.<br>303 18.                                       | 301 18.                                | 299 18.                      | 18.                    | 295 18.                                 | 293 18.<br>.294 18.   | 292 18.  | 289 18.<br>290 18.                                       | 18.                    | 18.                    |
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| 351 18 0.6 51<br>352 18 0.6 51<br>353 18 0.6 51<br>354 18 0.6 51<br>355 18 0.6 51<br>355 18 0.6 51  | 347     18     0.6     49       348     18     0.6     50       349     18     0.6     51       350     18     0.6     51 | 18 0.6 49<br>18 0.6 49<br>18 0.6 49                                  | 18 0.6 49<br>18 0.6 49<br>18 0.6 49                                  | 18 0.6 49<br>18 0.6 49                       | 18 0.6 45<br>18 0.6 49<br>18 0.6 49                                 | 334 18 0.6 45<br>335 18 0.6 45                   | 332 18 0.6 45<br>333 18 0.6 45                       | 18 0.6 45<br>18 0.6 45                             | $\begin{array}{cccccccccccccccccccccccccccccccccccc$    | 326 18 0.6 39<br>327 18 0.6 43                       | 324 18 0.6 36<br>325 18 0.6 37                 | 321 18 0.6 36<br>322 18 0.6 36<br>323 19 0.6 36                               | 319 18 0.6 35<br>320 18 0.6 35                     | 316 18 0.6 26<br>317 18 0.6 26 | 314 18 0.6 21<br>315 18 0.6 21   | 312 18.2 0.7 60<br>313 18.2 0.7 60                          | 310 18.2 0.7 60 6<br>311 18.2 0.7 60 9                                      | 308. 18.2 0.7 58 9<br>309 18.2 0.7 60 6                  | 306 18.2 0.7 57 5<br>307 18.2 0.7 58 6   | 304 18.2 0.7 56 6<br>305 18.2 0.7 56 9                                | 302 18.2 0.7 56 6<br>303 18.2 0.7 56 6                   | 300 18.2 0.7 54 9<br>301 18.2 0.7 56 6 | 299 18.2 0.7 54 6            | 18.2 0.7 51 13         | 295 18.2 0.7 51 9<br>296 18.2 0.7 51 10 | 293 18.2 0.7 51 6<br>.294 18.2 0.7 51 9                     | 291 18.2 0.7 51 6<br>292 18.2 0.7 51 6                   | 289 18.2 0.7 51 6<br>290 18.2 0.7 51 6                   | 18.2 0.7 51 6          | 18.2 0.7 51 6          |
| 351 18 0.6 51<br>352 18 0.6 51<br>353 18 0.6 51<br>354 18 0.6 51<br>355 18 0.6 51<br>355 18 0.6 51  | 347     18     0.6     49       348     18     0.6     50       349     18     0.6     51       350     18     0.6     51 | 18 0.6 49<br>18 0.6 49<br>18 0.6 49                                  | 18 0.6 49<br>18 0.6 49<br>18 0.6 49                                  | 18 0.6 49<br>18 0.6 49                       | 18 0.6 45<br>18 0.6 49<br>18 0.6 49                                 | 334 18 0.6 45<br>335 18 0.6 45                   | 332 18 0.6 45<br>333 18 0.6 45                       | 18 0.6 45<br>18 0.6 45                             | $\begin{array}{cccccccccccccccccccccccccccccccccccc$    | 326 18 0.6 39<br>327 18 0.6 43                       | 324 18 0.6 36<br>325 18 0.6 37                 | 321 18 0.6 36<br>322 18 0.6 36<br>323 19 0.6 36                               | 319 18 0.6 35<br>320 18 0.6 35                     | 316 18 0.6 26<br>317 18 0.6 26 | 314 18 0.6 21<br>315 18 0.6 21   | 312 18.2 0.7 60<br>313 18.2 0.7 60                          | 310 18.2 0.7 60 6 AR097345<br>311 18.2 0.7 60 9 HUMSAU3A1                   | 308. 18.2 0.7 58 9<br>309 18.2 0.7 60 6                  | 306 18.2 0.7 57 5<br>307 18.2 0.7 58 6   | 304 18.2 0.7 56 6<br>305 18.2 0.7 56 9                                | 302 18.2 0.7 56 6<br>303 18.2 0.7 56 6                   | 300 18.2 0.7 54 9<br>301 18.2 0.7 56 6 | 299 18.2 0.7 54 6            | 18.2 0.7 51 13         | 295 18.2 0.7 51 9 HUMCTVARA             | .293  | 291 18.2 0.7 51 6<br>292 18.2 0.7 51 6                   | 289 18.2 0.7 51 6<br>290 18.2 0.7 51 6                   | 18.2 0.7 51 6          | 18.2 0.7 51 6          |
| 351 18 0.6 51<br>352 18 0.6 51<br>353 18 0.6 51<br>354 18 0.6 51<br>355 18 0.6 51<br>356 18 0.6 51  | 347     18     0.6     49       348     18     0.6     50       349     18     0.6     51       350     18     0.6     51 | 18 0.6 49<br>18 0.6 49<br>18 0.6 49                                  | 18 0.6 49<br>18 0.6 49<br>18 0.6 49                                  | 18 0.6 49<br>18 0.6 49                       | 18 0.6 45<br>18 0.6 49<br>18 0.6 49                                 | 334 18 0.6 45<br>335 18 0.6 45                   | 332 18 0.6 45<br>333 18 0.6 45                       | 18 0.6 45<br>18 0.6 45                             | $\begin{array}{cccccccccccccccccccccccccccccccccccc$    | 326 18 0.6 39<br>327 18 0.6 43                       | 324 18 0.6 36<br>325 18 0.6 37                 | 321 18 0.6 36<br>322 18 0.6 36<br>323 19 0.6 36                               | 319 18 0.6 35<br>320 18 0.6 35                     | 316 18 0.6 26<br>317 18 0.6 26 | 314 18 0.6 21<br>315 18 0.6 21   | 312 18.2 0.7 60<br>313 18.2 0.7 60                          | 310 18.2 0.7 60 6 AR097345<br>311 18.2 0.7 60 9 HUMSAU3A1                   | 308. 18.2 0.7 58 9<br>309 18.2 0.7 60 6                  | 306 18.2 0.7 57 5<br>307 18.2 0.7 58 6   | 304 18.2 0.7 56 6<br>305 18.2 0.7 56 9                                | 302 18.2 0.7 56 6<br>303 18.2 0.7 56 6                   | 300 18.2 0.7 54 9<br>301 18.2 0.7 56 6 | 299 18.2 0.7 54 6            | 18.2 0.7 51 13         | 295 18.2 0.7 51 9 HUMCTVARA             | 293 18.2 0.7 51 6<br>.294 18.2 0.7 51 9                     | 291 18.2 0.7 51 6<br>292 18.2 0.7 51 6                   | 289 18.2 0.7 51<br>290 18.2 0.7 51                       | 18.2 0.7 51 6          | 18.2 0.7 51 6          |
| 351 18 0.6 51 6 AXI15589<br>352 18 0.6 51 6 AXI180013<br>353 18 0.6 51 6 AXI18101<br>354 18 0.6 51 6 AXI59785<br>355 18 0.6 51 6 AXI59937<br>356 18 0.6 51 6 AXI59937 | 347 18 0.6 49 6 AX077279<br>348 18 0.6 50 6 AX19802<br>349 18 0.6 51 6 AX7490<br>350 18 0.6 51 6 AR112992                 | 18 0.6 49 6 AR087689<br>18 0.6 49 6 AR088196<br>18 0.6 49 6 AR094049 | 18 0.6 49 6 ARO58370<br>18 0.6 49 6 ARO88739<br>18 0.6 49 6 ARO84881 | 18 0.6 49 6 AR042490<br>18 0.6 49 6 AR055555 | 18 0.6 45 6 1910/98<br>18 0.6 49 6 AR013876<br>18 0.6 49 6 AR033830 | 334 18 0.6 45 6 129424<br>335 18 0.6 45 6 125019 | 332 18 0.6 45 6 AX306498<br>333 18 0.6 45 6 AX382512 | 18  0.6  45  6 AR032684<br>18  0.6  45  6 AR209348 | 18 0.6 43 6 ARO34560<br>18 0.6 43 6 I89250              | 326 18 0.6 39 9 S77756<br>327 18 0.6 43 6 ARO04648 . | 324 18 0.6 36 115726<br>325 18 0.6 37 6 121496 | 321 18 0.6 36 ARO30428<br>322 18 0.6 36 6 ARO30446<br>323 18 0.6 36 6 ARO3045 | 319 18 0.6 35 6 AR063739<br>320 18 0.6 35 6 I30102 | 310                            | 314 18 0.6 21 6 AX096028<br>315 18 0.6 21 6 AX096030                             | 312 18.2 0.7 60 9 HUMSAU3A35<br>313 18.2 0.7 60 12 SYNIGMAB | 310 18.2 0.7 60 6 'AR097345<br>311 18.2 0.7 60 9 HUMSAU3A12                 | 308. 18.2 0.7 58 9 MTHSLAS98<br>309 18.2 0.7 60 6 A76630 | 306 18.2 0.7 57 5 SMOACACB<br>307 18.2 0.7 58 6 AX148297                       | 304 18.2 0.7 56 6 AX184575<br>305 18.2 0.7 56 9 S63111                | 302 18.2 0.7 56 6 AX089727<br>303 18.2 0.7 56 6 AX090182 | 301 18.2 0.7 56 6 AX089726             | 299 18.2 0.7 54 6 AX254831 . | 18.2 0.7 51 13 \$76572 | 295 18.2 0.7 51 9 HUMCTVARA             | 293 18.2 0.7 51 6 AX204366<br>.294 18.2 0.7 51 9 HUMD6C04M3 | 291 18.2 0.7 51 6 AX165676<br>292 18.2 0.7 51 6 AX199233 | 289 18.2 0.7 51 6 AXI65384<br>290 18.2 0.7 51 6 AXI65613 | 18.2 0.7 51 6 AX163333 | 18.2 0.7 51 6 AX160457 |
| 351 18 0.6 51 6 AXI15589<br>352 18 0.6 51 6 AXI180013<br>353 18 0.6 51 6 AXI18101<br>354 18 0.6 51 6 AXI59785<br>355 18 0.6 51 6 AXI59937<br>356 18 0.6 51 6 AXI59937 | 347 18 0.6 49 6 AX077279<br>348 18 0.6 50 6 AX19802<br>349 18 0.6 51 6 AX7490<br>350 18 0.6 51 6 AR112992                 | 18 0.6 49 6 AR087689<br>18 0.6 49 6 AR088196<br>18 0.6 49 6 AR094049 | 18 0.6 49 6 ARO58370<br>18 0.6 49 6 ARO88739<br>18 0.6 49 6 ARO84881 | 18 0.6 49 6 AR042490<br>18 0.6 49 6 AR055555 | 18 0.6 45 6 1910/98<br>18 0.6 49 6 AR013876<br>18 0.6 49 6 AR033830 | 334 18 0.6 45 6 129424<br>335 18 0.6 45 6 125019 | 332 18 0.6 45 6 AX306498<br>333 18 0.6 45 6 AX382512 | 18  0.6  45  6 AR032684<br>18  0.6  45  6 AR209348 | 18 0.6 43 6 ARO34560<br>18 0.6 43 6 I89250              | 326 18 0.6 39 9 S77756<br>327 18 0.6 43 6 ARO04648 . | 324 18 0.6 36 115726<br>325 18 0.6 37 6 121496 | 321 18 0.6 36 ARO30428<br>322 18 0.6 36 6 ARO30446<br>323 18 0.6 36 6 ARO3045 | 319 18 0.6 35 6 AR063739<br>320 18 0.6 35 6 I30102 | 310                            | 314 18 0.6 21 6 AX096028<br>315 18 0.6 21 6 AX096030                             | 312 18.2 0.7 60 9 HUMSAU3A35<br>313 18.2 0.7 60 12 SYNIGMAB | 310 18.2 0.7 60 6 'AR097345<br>311 18.2 0.7 60 9 HUMSAU3A12                 | 308. 18.2 0.7 58 9 MTHSLAS98<br>309 18.2 0.7 60 6 A76630 | 306 18.2 0.7 57 5 SMOACACB<br>307 18.2 0.7 58 6 AX148297                       | 304 18.2 0.7 56 6 AX184575<br>305 18.2 0.7 56 9 S63111                | 302 18.2 0.7 56 6 AX089727<br>303 18.2 0.7 56 6 AX090182 | 301 18.2 0.7 56 6 AX089726             | 299 18.2 0.7 54 6 AX254831 . | 18.2 0.7 51 13 \$76572 | 295 18.2 0.7 51 9 HUMCTVARA             | 293 18.2 0.7 51 6 AX204366<br>.294 18.2 0.7 51 9 HUMD6C04M3 | 291 18.2 0.7 51 6 AX165676<br>292 18.2 0.7 51 6 AX199233 | 289 18.2 0.7 51 6 AXI65384<br>290 18.2 0.7 51 6 AXI65613 | 18.2 0.7 51 6 AX163333 | 18.2 0.7 51 6 AX160457 |
| 351 18 0.6 51<br>352 18 0.6 51<br>353 18 0.6 51<br>354 18 0.6 51<br>355 18 0.6 51<br>355 18 0.6 51  | 347 18 0.6 49 6 AX077279<br>348 18 0.6 50 6 AX19802<br>349 18 0.6 51 6 AX7490<br>350 18 0.6 51 6 AR112992                 | 18 0.6 49 6 AR087689<br>18 0.6 49 6 AR088196<br>18 0.6 49 6 AR094049 | 18 0.6 49 6 ARO58370<br>18 0.6 49 6 ARO88739<br>18 0.6 49 6 ARO84881 | 18 0.6 49 6 AR042490<br>18 0.6 49 6 AR055555 | 18 0.6 45 6 1910/98<br>18 0.6 49 6 AR013876<br>18 0.6 49 6 AR033830 | 334 18 0.6 45 6 129424<br>335 18 0.6 45 6 125019 | 332 18 0.6 45 6 AX306498<br>333 18 0.6 45 6 AX382512 | 18  0.6  45  6 AR032684<br>18  0.6  45  6 AR209348 | 18 0.6 43 6 ARO34560<br>18 0.6 43 6 I89250              | 326 18 0.6 39 9 S77756<br>327 18 0.6 43 6 ARO04648 . | 324 18 0.6 36 115726<br>325 18 0.6 37 6 121496 | 321 18 0.6 36 ARO30428<br>322 18 0.6 36 6 ARO30446<br>323 18 0.6 36 6 ARO3045 | 319 18 0.6 35 6 AR063739<br>320 18 0.6 35 6 I30102 | 310                            | 314 18 0.6 21 6 AX096028<br>315 18 0.6 21 6 AX096030                             | 312 18.2 0.7 60 9 HUMSAU3A35<br>313 18.2 0.7 60 12 SYNIGMAB | 310 18.2 0.7 60 6 AR097345 AR097345; 311 18.2 0.7 60 9 HUMSAU3A12 D49596 Hu | 308. 18.2 0.7 58 9 MTHSLAS98<br>309 18.2 0.7 60 6 A76630 | 306 18.2 0.7 57 5 SMOACACB M19425 Ra.<br>307 18.2 0.7 58 6 AX148297 AX148297 : | 304 18.2 0.7 56 6 AX184575 AX184575 305 18.2 0.7 56 9 S63111 S63111 T | 302 18.2 0.7 56 6 AX089727<br>303 18.2 0.7 56 6 AX090182 | 301 18.2 0.7 56 6 AX089726             | 299 18.2 0.7 54 6 AX254831 . | 18.2 0.7 51 13 \$76572 | 295 18.2 0.7 51 9 HUMCTVARA             | 293 18.2 0.7 51 6 AX204366<br>.294 18.2 0.7 51 9 HUMD6C04M3 | 291 18.2 0.7 51 6 AX165676<br>292 18.2 0.7 51 6 AX199233 | 289 18.2 0.7 51 6 AXI65384<br>290 18.2 0.7 51 6 AXI65613 | 18.2 0.7 51 6 AX163333 | 18.2 0.7 51 6 AX160457 |

|          |   |         |               |            |         |          |       |         |         |       |               |       |               |               |               |                        |                  |           |            |         |         |         |       |         |         |               |                 |       |            |           |         |         |         |              |         |               |                |         |       |       |          |          |         |         |         |            |        |         |         |       |         |   |                                       |         | ٠     |          |       |            |         |         |    |
|----------|---|---------|---------------|------------|---------|----------|-------|---------|---------|-------|---------------|-------|---------------|---------------|---------------|------------------------|------------------|-----------|------------|---------|---------|---------|-------|---------|---------|---------------|-----------------|-------|------------|-----------|---------|---------|---------|--------------|---------|---------------|----------------|---------|-------|-------|----------|----------|---------|---------|---------|------------|--------|---------|---------|-------|---------|---|---------------------------------------|---------|-------|----------|-------|------------|---------|---------|----|
| c 429    | C 427   | 425     |               | C 423      | C 421   | 420      | 419   | 418     |         | c 416 | 415           | 414   |               | c 412         |               |                        | 0 459            |           | 400        | 0 405   |         | 403     | 402   | 401     | 400     |               |                 | 707   |            |           |         | c 352   |         | c 390        |         |               | C 387          | 0 385   |       |       | c 382    |          | 379     | 378     | 377     |            | 0 375  |         | c 372   |       |         |   | 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 |         | 365   | 364      | 363   | 362        |         | 0 359   |    |
|          | <u>, , , , , , , , , , , , , , , , , , , </u> | سز د    | ₽,            |            | د       |          | 1     | ш       |         | ב     |               |       | . د           | <u>.</u> ,    | _             | - ,                    | <u>.</u> ,       |           | ـر د       | ــو د   |         |         |       | _       |         | <b>–</b>      | <b>⊸</b> 1      |       |            | ــر د     | . ــ    |         | <b></b> | 1            |         | _             |                |         |       |       |          |          |         |         |         |            |        |         |         |       |         |   |                                       |         |       |          |       |            |         |         |    |
|          | 7.8   |         |               |            |         |          |       |         |         |       |               |       |               |               |               |                        |                  |           |            |         |         |         |       |         |         |               |                 |       |            |           |         |         |         |              |         |               | 18             | 1 8     | 18    | 18    | 18       | 1 A      | 18      | 18      | 18      | 18         | 0 a    | 18      | 18      | 18    | 18      | 18                                      | 3 5                                   | ο α     | 18    | 18       | 18    | 180        | 0 20    | 2 2     | دُ |
|          | 000   |         |               |            |         |          |       | •       |         |       |               |       |               |               |               |                        |                  |           |            |         |         |         |       |         |         |               |                 |       |            |           |         |         |         |              |         |               |                |         |       |       |          |          |         |         |         |            |        |         |         |       |         |   |                                       |         |       |          |       |            |         |         |    |
| 50       | 444   | 4 9     | 49            | 4 4<br>9 9 | 49      | 48       | 48    | 48      | 48      | 48    | 4-<br>00      | 47    | 47            | 47            | 47            | 47                     | 47               | 7 0       | 4 4        | 4.0     | 4.5     | 45      | 43    | 43      | 43      | 4.            | 40              | ມູດ   | . u        | υ<br>0 00 | 38      | 38      | 31      | 30           | 30      | 28            | n o            | 60      | 60    | 60    | 60       | n 0      | 00      | 59      | 59      | ن ر<br>د د | n 0    | י ט     | 55      | 54    | 54.     | 5.                                      | л U                                   | 5.2     | 5 2   | 51       | 51    | <u>ا ٦</u> | 5 J     | 5       | 1  |
|          | 0 0 0<br>2 2 2                                |         |               |            |         | 0        |       |         |         |       |               |       |               |               |               |                        |                  |           | J          |         |         |         |       |         |         |               |                 |       |            |           |         |         |         |              |         | 4             | ~ ~            | ١.      |       |       |          |          |         |         |         | _          | J      |         |         |       |         |   |                                       |         |       | 0        |       |            |         |         |    |
| X161238  | AX356711<br>AX356711                          | X259764 | R210917       | R210916    | R153115 | MUSIGHZB | 40904 | X222045 | X081679 | 78687 | 78686         | 41063 | 30213         | X194718       | X194680       | R088352                | R036885          | 51 NGALAL | 14204      | X236658 | R147495 | R125138 | 38119 | R077198 | R017793 | R011922       | X456405         | 52436 | X2/3588    | R148761   | R045396 | R045384 | X249324 | R064922      | R000099 | R092331       | SHITRANS       | NS019CB | 20529 | 09144 | 05516    | X02/9080 | R117072 | D001723 | D001710 | X148295    | EBSAT4 | X485745 | R088559 | 81114 | X303573 | X076553                                 | AR205535                              | R083330 | 50699 | MMU90451 | 61342 | X204092    | X199613 | X199611 |    |
|          |   |         |               |            |         |          |       |         |         |       |               |       |               |               |               |                        |                  |           |            |         |         |         |       |         |         |               |                 |       |            |           |         |         |         |              |         | 2             | .T.A           |         |       |       |          |          |         |         |         |            |        |         |         |       |         |   |                                       |         |       | •        |       |            |         |         |    |
|          |   |         |               |            |         |          |       |         |         |       |               |       |               |               |               |                        |                  |           |            |         |         |         |       |         |         |               |                 |       |            |           |         |         |         |              |         |               |                |         |       |       |          |          |         |         |         |            |        |         |         |       |         |   |                                       |         |       |          |       |            |         |         |    |
| AX<br>AX | A X   | AX      | AR            | A R        | AR      | ×        | E4    | ΑX      | AX      | Α7    | A7            | I4    | 133           | AX            | AX.           | A                      | מ מ              | ,<br>,    | K F        | AX      | AR      | AR      | Ε3    | AR      | AR      | AR.           | P F             | 7. U  | τAX        | AR        | AR      | AR      | AX      | AR           | AR.     | Þ.            | X D            | ΑL      | 12    | 10    | IO.      | AX       | AR      | BD      | BD :::  | × ×        | ×      | AX      | AR      | 88    | A S     | A S                                     | A A                                   | AR      | A5    | U        | E 6   | A A        | . AX    | AX      | ,  |
| 16123    | AX356/10 Sequence<br>AX356711 Sequence        | 25976   | 21091         | 210G1      | 15311   | 22398    | 0904  | 22204   | 08167   | 8687  | 8686          | 1063  | 0213          | 79471         | 19468         | 08835                  | 20000            | 102507    | 4204       | 23665   | 14749   | 12513   | 8119  | 07719   | 01779   | 01192         | 72727           | 2440  | 2/358      | 14876     | 04539   | 04538   | 24932   | 06492        | 500000  | 20073         | 27001          | 11145   | 0529  | 9144  | 5516     | 707      | 11707   | 00172   | 00171   | 14820      | 9458   | 48574   | 08855   | 1114  | 30357   | 0765                                    | 20557                                 | 08333   | 9699  | 190451   | 1342  | 20405      | 19961   | 19961   |    |
| a Sec    | 1 Sec   | 4 Sec   | 7 Sec         | 20.0       | 5 Sec   | Mous     | Humar | 5 Sec   | 9 Sec   | Seque | Segne         | Seque | Segue         | 200           | Sec           | Sec                    | ο<br>N<br>O<br>O | 0 0 0 0   | PCR 1      | Sec Sec | 5 Sec   | 8 Sec   | DNA ( | 8 Sec   | 3 Sec   | Sec           | 200             | seque | Sec Sec    | 1 Sec     | 6 Sec   | 4 Sec   | 4 Sec   | 22 Sec       | 99 Sec  | 200           | 28053 Sequence | 9 Bo    | Sequi | Sequi | Segu     | 000      | 2 Sec   | 3 Me    | 000     | 5 25       | C.ap   | 5 Se    | 9 Se    | Homo  | Sec     | 200                                     | 5 C                                   | Se Se   | Sequ  | Mus      | dord  | 000        | Se Se   | 11 Se   |    |
| gnenc    | quenc   | quenc   | duenc         | quenc      | quenc   | se Ig    | nized | quenc   | nenc    | ence  | ance<br>once  | ence  | ance<br>Arene | oudill.       | Jueno         | Judiik                 | Judin.           | foral     | brime      | quenc   | quenc   | quenc   | elong | quenc   | quenc   | duenc         | and and         | ence  | quenc      | quenc     | quenc   | quenc   | quenc   | quenc        | guenc   | dilent        | equer          | tryti   | ence  | ence  | ence     | quenc    | quenc   | thod    | i sodu  | apter      | eLla   | quenc   | quenc   | sapi  | duenc   | dieno                                   | quenc                                 | quenc   | ence  | musc     | e foi | queno      | quenc   | quenc   |    |
| m m      | . n n   | n       | no n          |            | O       | зe       | שנ    | (D      | ው ‹     | 4     | ، بد          | O) (  | ט עכ          | D (           | D C           | D A                    | Dα               | 5111      | ; H        | ው       | Ф       | Ð       | a÷    | ው 1     | O (     | D (           | D F             | 10    | 3 0        | O         | Ф       | Ð       | Œ       | ወ (          | O (     | י ט           | 000            | S       | 72    | 4     | 1) 4     | o (0     | O       | fo      | ţ. (    | יט<br>כ    | Sa     | Ф       | O       | en    | D (I    | DC                                      | D (D                                  | Ð       | 38    | ulc      | α "   | Ď Œ        | Ō       | Ø       |    |
|          |   | . ,     |               |            |         | -        |       |         |         |       |               |       |               |               |               |                        |                  |           |            |         |         |         | _     |         |         |               |                 |       |            |           |         |         |         |              |         |               |                |         | _     |       |          |          | _       |         |         |            |        |         |         |       |         |   |                                       |         |       |          |       |            |         |         |    |
|          | c 499   | 4.0     | 4 4           | 4 4        | 5       | 49       | 4.9   | 4       | 4       | C 489 | 4 4           | m d   | 4 4 6 0 0     |               | 4 4           | 4 4                    | 4 4              | 4.4       | C 479      | 4.      | 47      | c 476   | 475   | 474     | 4.      | c 472         | 471             | 4.    | 0 468      | 4.        | 466     | 46      | C 464   | 4.           | c 462   | 460           | 4.4            | 4       | c 457 | 4.    | 4 4      | 4.       | c 452   | 4       | 4       | 448        |        | ~       | 4.      |       | 4 4     | 4 4                                     | 440                                   |         |       |          |       | 4          | c 433   | 4       |    |
|          | 17.6<br>17.6                                  |         |               |            |         |          |       |         |         |       |               |       |               | 17.6          |               | 17.6                   |                  |           | 17.8       |         |         |         |       |         |         |               |                 | 17.8  | 17.8       |           |         |         |         |              |         |               | 17.8           |         |       |       | 17.8     |          | 17.8    |         |         | 17 g       |        |         |         |       |         |   | 17.E                                  |         |       |          | 17.6  |            |         |         |    |
|          | 0.0   | 0       | 00            | 0.0        | 0.      | 0.       | 0.    | 0       | 0 9     | 0 9   | <b>&gt;</b> ( | 0 9   | <b>.</b>      | <b>&gt;</b> : | <b>&gt;</b> ( | > <                    | <b>&gt;</b>      |           |            | 0       | 0.      | 0.      | 0     | 0       | 0       | <b>&gt;</b> • | <b>5</b>        | 9.5   | 0          | 0         | 0       | 0       | 0       | 0 9          | 0 0     | <b>&gt;</b> < |                | 0       | 0     | 0     | 00       | 0        | 0       | 0       | 0 (     | ٥ د        | 0      | 0       | 0       | 0     | ٥ د     | <b>&gt;</b>                             | 00                                    | . 0     | 0     | 0        | 0 0   | -<br>-     | . 0     | . 0     | ,  |
|          | თთ  |         |               |            |         |          |       |         |         |       |               |       |               |               | •             |                        |                  |           |            |         |         |         |       |         |         |               |                 |       |            |           |         |         |         |              |         |               |                |         |       |       |          |          |         |         |         |            |        |         |         |       |         |   |                                       |         |       |          |       |            |         |         |    |
|          | 41 6  |         |               |            |         |          |       |         |         |       |               |       |               |               |               |                        |                  |           |            |         | _       | _       | _     | _       | _       |               |                 |       |            |           | _       | _       |         |              | _       |               |                | _       | _     |       |          |          |         | _       | _       |            |        | _       | _       | _     |         |   |                                       |         | _     | _        | ٠.    |            |         | _       |    |
|          | 6 AR03  |         |               |            | 11      | EL       | AR:   | AR      | AR      | A1    | 18            | AR    | A A           | N D           | N C           | 0 2                    | ) L              | AK        | HU         | HS      | .E2     | AX      | AX    | AR      | A :     | 21            | 2 2             | A     | AR         | 12        | 5 17    | AR      | AF      | M (          | 16      | 7. L          | I AR           | AR      | 5 AR  | AR:   | AX183597 | . A      | 14 A    | AX      | AX:     | . O        | AX     | AX      | ĀX      | AX :  | A 2     | )                                       | A A                                   | AX      | AX    | AX       | ) N   | , A        | AX      | AX      |    |
|          | 03677   | 8       | 57            | 5123       | 1602    | 4547     | 13343 | 05843   | 05842   | 7744  | 1283          | 18230 | 1348¢         | 31734         | 22526         | 55576                  | 3707:            | 1737      | HUMSAU3A58 | EZRIN   | 2154    | 38250   | 36597 | 09734   | 9222    | 2027          | 0440            | 4/432 | 05220      | 8251      | 5358    | 12768   | 06741   | 7060W        | 21.0    | 7160          | 20306          | 13548   | 05916 | 0412  | 18350    | F233     | F233    | 3222    | 20430   | 755UC      | 16554  | 16524   | 1650    | 16498 | 1627    | 7 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 | 1630                                  | 1622    | 1585  | 1585     | 1577  | 1559       | 1165    | 11628   |    |
|          | ώ α   | ŭ       | ŏ             |            |         |          | 9     | õ       | 9       |       | ì             | 3     | تَ مَ         | . 6           | 5 6           | " č                    | ñ                | Č         | 8A58       | 1G16    |         | )7      | 17    | 4       |         |               |                 | 2.2   | ; <u>:</u> | •         |         | 39      | 6       | 86           |         |               | 2              | S S     | 59    | 24    | 7 6      | 285      | 081     | 20      | 0       | 2 6        | 2      | 9       | 59      | 8.5   | 72      | 71                                      | 2 6                                   | 3 5     | 30    | 29       | 2.2   | 5 1        | 29      | 39      |    |
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|          |   |         |               |            |         |          |       |         |         |       |               |       |               |               |               |                        |                  |           |            |         |         |         |       |         |         |               |                 |       |            |           |         |         |         |              |         |               |                |         |       |       |          |          |         |         |         |            |        |         |         |       |         |   |                                       |         |       |          |       |            |         |         |    |
|          | AR036778 Sequence<br>BD012559 Method of       | AR20    | AX07          | 1351       | 1116    | E145     | AR13  | AR05    | AROS    | A177  | 1812          | AR18  | A R 1 3       | 7770          | AXXX          | 2075                   | 7777             | AKIC      | D496       | AF05    | E221    | AX38    | AX36  | AROS    | A292    | 19C           | ALOS            | AX4   | AROS       | 1282      | I753    | AR12    | AF06    | )60<br>260 T | 165E    | LVCI          | AR20           | AR1     | ARO:  | ARO.  | AX18     | AF       | AF      | AX3     | AX20    | AXA        | AX16   | AX16    | AX16    | AX16  | AXI     | 271                                     | AXI                                   | AX16    | AX15  | AX1      | AXI   | AXL        | AX1     | AX1     |    |
|          | 2559  | )8355   | 6796          | 123 S      | 02 S    | 547 D    | 3439  | 58430   | 8429    | 744 N | 2000          | 12309 | 11243         | 72/2          | 35360         | מים מוני               | 77775            | 74/90     | 542 H      | 53446   | 154 Y   | 32507   | 55977 | 77344   | 222 0   | 7 C E C       | 721 N           | 14322 | 52201      | 251 S     | 358 S   | 27689   | 57416   | )498         | 2100    | 2000          | 03062          | 35485   | 59169 | 11224 | 33597    | 23318    | 23318   | 22220   | 04300   | 7//59      | 55542  | 55249   | 55059   | 54986 | 57777   | 7777                                    | 53002                                 | 52213   | 58530 | 58529    | 57742 | 57503<br>1 | 16529   | 16289   |    |
|          | Sequ<br>Meth                                  | Sequ    | nbəs<br>Ianha | equer      | equer   | NA pr    | Sequ  | Sequ    | Seal    | uclec | Sea of        | reds. | Sedi          | 500           | Sec           | - 72 64                | ianba            | Sequ      | uman       | Home    | east    | Sequ    | Sequ  | Sequ    | ligor   | וויט ויסל     | Tigo            | Sequ  | Sequ       | equei     | equer   | Sequ    | Home    | Mus musculu  | edile   | edue          | Seq            | Seq     | Sequ  | Sequ  | Sec      | 5 Hui    | 0 Hui   | Sequ    | Segu    | Sed        | Seq    | Sequ    | Sequ    | Sequ  | Sec     | 500                                     | Seq                                   | Sequ    | Sequ  | Sequ     | Sed   | seq        | Seq     | Seq     |    |
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Nucleic acids containing single nucleotide
methods of use thereof
                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                              /note="single nucleotide polymorphism
Accession number cg43297259"
a 14 c 12 g 15 t
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Nucleic acids containing single nucleotide
methods of use thereof
Patent: WO 0140521-A 4583 07-JUN-2001;
Curagen Corporation (US)
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AX161256
AX161256.1 GI:14542587
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Eukaryota; M
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79.6%;
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AX161706/c
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AX163066
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              1687 CATTTGTATGTGCTGAGTGTGGACGAGGCTTTAATGATAAGTCCACCCTC 1736
 51
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AX161706
AX161706.1
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AX163066
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Curagen
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1 (bases 1 to 51)
Shimkets, R.A. and Leach, M.
Nucleic acids containing single nucleotide polymorphisms and methods of use thereof
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35; Conserv
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Patent: WO 0140521-A 6394 07-JUN-2001;
Curagen Corporation (US)
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                                                                                                                                                                                                                                                                                                                                         Homo sapiens
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                                                                  Similarity
CTTATGAATGTACAGAGTGTGGGAAAGCCTTCAAACATAGTTCCACCCTC
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Corporation (US)
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                                                                                                                   /note="2 of 2 allelic variants (5033 is other entry) Accession number cg43981141" a 10 c 13 g 15 t .
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                                                                                                                                                              /organism="Homo sapiens"
/db_xref="taxon:9606"
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/db_xref="taxon:9606"
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Catarrhini; Hominidae; Homo.
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RESULT 7

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AX199456/c
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Ax382519
Sequence 49
Ax382519
Ax382519.1
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Sequence
AX199456
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Zinc finger domain recognition code a
Patent: WO 0208286-A 36 31-JAN-2002;
Syngenta Participations AG (CH)
Syngenta Participations AG (CH)
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Sequence 36 from Patent W00208286.
AX382506
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                                                                                                                                                     Similarity 71.4
35; Conservative
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386 from Patent WO0151670
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/db_xref="taxon:9606"
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/db_xref="taxon:32630"
/note="Partial zinc finger doma
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                       56 bp
from Patent WO0208286
  GI:19577270
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Pred. No. 1.8e+04;
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RESULT 11
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Sequence 5033 from F
AX161705
AX161705.1 GI:14543
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 Eukaryota; Metazoa; Mammalia; Eutheria;
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/db_xref="taxon:9606"
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6393 from Patent WO0140521.
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/db_xxef="taxon:32630"
/db_xzef="Partial zinc finger domain
/note="Partial zinc finger domain
10 c 19 g 12 t
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71.4%;
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 Chordata;
Primates;
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Pred. No. 3.2e
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Craniata; Ve Catarrhini;
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            Vertebrata;
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            Euteleostomi;
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AX382515
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KEYWORDS
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AX382510
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Shinkets, R.A. and Leach, M.
Nucleic acids containing single nucleotide
methods of use thereof
Patent: WO 0140521-A 5033 07-JUN-2001;
Curagen Corporation (US)
Curagen Corporation (US)
Sera,T.

Zinc finger domain recognition code.
Patent: WO 0208286-A 40 31-AAN-2002;
Syngenta Participations AG (CH)
Syngenta Participations AG (CH)
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Sequence
AX382510
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AX382515
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/db_xref="taxon:9606"
26
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/db_xref="taxon:32630"
/note="Partial zinc finger domain
9 c 19 g 14 t
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AX199103
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AX199104
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AX199103
AX199103.1
                                                   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 51)

Shimkets,R.A. and Leach,M.D.

Nucleic acids containing single nucleotide polymorphisms and methods of use thereof
Patent: WO 0151670-A 34 19-JUL-2001;
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Sequence 34 from Patent
AX199104
AX199104.1 GI:15389448
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                                             Curagen
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1 (bases 1 to 51)
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31; Conservative
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/organism="synthetic construct"
/db_xref="taxon:32630"
/note="Partial zinc finger domain oligomer.
/note="Partial zinc finger domain oligomer.
a 9 c 16 g 10 t 8 others
                                                                                                                                                                                                                                                                                                                                                                                             /organism="Homo sapiens"
/db_xref="taxon:9606"
26
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10 c
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Accession number cg43926631"
10 c 16 g 12 t
 /db_xref="taxon:9606"
           /organism="Homo sapiens"
                                 Location/Qualifiers
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Pred. No. 9.9e+04;
0; Mismatches 14;
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AX163385/c
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1211 CTTTAGCCTGAAGTCAAACCTCATTACCCACCAGAGGGCGCACACTGGGGAGAAG 1265
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                                                                                                                                                                                                                                                                          Sequence 67
AX163385
AX163385.1
                                                                                                Mammalia: Eutheria; Primates; Catarrhini; H
1 (bases 1 to 51)
Shimkets, R.A. and Leach, M.
Nucleic acids containing single nucleotide methods of use thereof
Patent: Wo 0140521-A 6713 07-JUN-2001;
Curagen Corporation/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 Similarity
36; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Zinc finger domain recognition code
Patent: WO 0208286-A 50 31-JAN-2002;
Syngenta Participations AG (CH)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            synthetic construct.
                                                                                                                                                                                                                                   Homo sapiens
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                                           /organism="Homo sapiens"
/db_xref="taxon:9606"
26
/note="1 of 2 allelic variants
Accession number cg43280136"
a 13 c 17 g 6 t
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Accession number cg43s
10 c 15 g
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/db_xref="taxon:32630"
/note="Partial zinc finger doma
14 c 15 g 17 t
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Pred. No. 1e+0
0; Mismatches
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Pred. No. 9.9e+04;
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AX382511/c
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KEYWORDS
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AX382511
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            synthetic construct. synthetic construct artificial sequences.
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Patent: WO 0140521-A 2987 07-JUN-2001;
Curagen Corporation (US)
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34; Conservative
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                                                                                         /organism="Homo sapiens"
/db_xref="taxon:9606"
26
                                                      /note="1 of 2 allelic variants (2988 Accession number cg42708452" 9 c 12 9 9 t
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/db_xref="taxon:32630"
/note="Partial zinc finger domain oligomer."
/note="partial zinc finger domain oligomer."
/note="partial zinc finger domain oligomer."
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Pred. No. 1.8e+05;
0; Mismatches 24;
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Pred. No. 1.8e+05;
0; Mismatches 17;
Score 23; DB 6; Le
Pred. No. 3.1e+05;
0; Mismatches 10;
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AX199545
AX199545.1
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Nucleic acids containing single nucle
methods of use thereof
Patent: WO 0151670-A 475 19-JUL-2001;
Curagen Corporation (US)
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32; Conserv
                                                                                                                                                                                                                   Wang, Z. and Xiao, W. A method for prognosing cancer and the Patent: Wo 0206327-A 34 24-JAN-2002; Northwestern University (US)
                                                                                                                                                                                                                                                                                                             synthetic construct. synthetic construct
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TCAGGATCCGCGCGTGCACCCGCGTTCCATCTGTCTTCTGAGACTT
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/db_xref="taxon:9606"
26
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/db_xref="taxon:32630"
/note="primer"
16 c 17 g 16 t
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Pred. No. 3.1e+05;
                                                         Score 23; DB 6;
Pred. No. 3.2e+05;
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WO0206327.
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AX382509/c
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                                                                                       synthetic synthetic
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29; Conser
          Zinc finger domain recognition code and Patent: WO 0208286-A 39 31-JAN-2002; Syngenta Participations AG (CH)
                                                                                                                                          Sequence 39
AX382509
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2 (bases 1 to 48)
Lichter,P., Bray,P., Ried,T., Dawid,I.B. and War
Clustering of C2-H2 zinc finger motif sequences
and fragile site regions of human chromosomes
Genomics 13 (4), 999-1007 (1992)
92372070
                                                                                                                               AX382509.1
                                                                                                                                                                   AX382509
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Bray, P., Lichter, P., Thiesen, H.J., War Characterization and mapping of human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota; Metazoa; Chordata; Mammalia; Eutheria; Primates;
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Eukaryota; Metazoa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DNA-binding protein; zinc Homo sapiens Placenta DNA
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                                                                           urtificial sequences.
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                                                                                                                                                                                                                                                                                                                                                 /translation="IHTGEKPFSCRECSKA"
15 c 17 g 6 t
                                                                                                                                                                                                                                                                                                                                                                      /product="DNA-binding protein"
/protein_id="AAA61327.1"
/db_xref="GI:340476"
Location/Qualifiers
                                                                                       construct.
                                                                                                                                                                                                                                                                                                                                                                                                                         /standard_name="zinc finger motif"
/note="putative"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /organism="Homo sapiens"
/db_xref="taxon:9606"
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/gene="ZNF"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /gene="ZNF"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              'tissue_type="Placenta"
                                                                                                                               GI:19577260
                                                                                                                                                       from Patent
                                                                                                                                                                                                                                                                                             0.8%;
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Pred. No. 4.7e+05;
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WO0208286.
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Catarrhini; Hominidae;
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man genes encoding zinc
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                                      uses thereof
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partial cds.
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AX382518/c
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Zinc finger domain recognition code and Patent: WO 0208286-A 48 31-JAN-2002; Syngenta Participations AG (CH)

Location/Qualifiers
                                                                                                              Mammalia: Eutheria; Primates; Catarrhini; H
1 (bases 1 to 51)
Shimkets, R.A. and Leach, M.
Nucleic acids containing single nucleotide
methods of use thereof
Patent: Wo 0140521-A 6714 07-JUN-2001;
Curagen Corporation (US)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     60 bp
Sequence 48 from Patent W00208286
AX382518
                                                                                                                                                                                                                                                                       Sequence 6714
AX163386
                                                                                                                                                                                                                    Homo sapiens
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synthetic construct
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                                                 /organism="Homo sapiens"
/db_xref="taxon:9606"
26
         /note="2 of 2 allelic variants (6713 Accession number cg43280136" 12 c 17 g 7 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /organism="synthetic construct"
//db_xxef="taxon:32630"
/note="Partial zinc finger domain
11 c 19 g 15 t
                                                                                                   Location/Qualifiers
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81.2%;
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                                                                                                                                                                                                                                                                                                                                                                                                                  0;
                                                                                                                                                                                                                                                                                                                                                                                                                 Score 22.4; DB 6;
Pred. No. 4.9e+05;
0; Mismatches 6;
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WO0140521.
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Query Match

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22.2;

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RESULT 28
AF062523
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AX115765/c
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ORGANISM
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AR036764/c
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                                                                                                                                                                                                                              AUTHORS
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AX115765
AX115765.1
                                                                                                    Similarity
27; Conserv
 AF062523
                                                                                                                                                                                                       Nucleic acid sequence detection by site in amplification reactions Patent: US 5800984-A 6 01-SEP-1998;
                                                                                                                                                                                                                                        l (bases 1 to 41)
Vary, C.P.H.
                                                                                                                                                                                                                                                                           Unknown
                                                                                                                                                                                                                                                                                      Unknown.
                                                                                                                                                                                                                                                                                                            AR036764.1
                                                                                                                                                                                                                                                                                                                       Sequence 6 from AR036764
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Genotyping reagents, kits and methods of use thereof Patent: WO 0129262-A 888 26-APR-2001; Orchid BioSciences, Inc. (US)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Picoult-Newburg, L. and Pohl, M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Eukaryota; Metazoa; Chordata; Mammalia; Eutheria; Primates; 1 (bases 1 to 51)
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                                                       GCGAGGGAAGCGCCCTGGAGAGAGAGAGAAAATG 409
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                                                                                                                                                           /organism="unknown"
16 c 7 g
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/db_xref="taxon:9606"
9 c 14 g 1
                                                                                                                                                                                             Location/Qualifiers
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70.7%;
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                                                                                                                                                                                                                                                                                                                                 patent US
                                                                                                   Score 21.6; DB 6;
Pred. No. 8.2e+05;
0; Mismatches 9
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0; Mismatches 18;
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AX159660/c
LOCUS
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              2723_GAGAGCTGTTCTTTTGCTCAATAAAATTCTTTTCTACCC
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33;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                369 CTGGAGCGAGGGGAAGCGCCCTGGAGAGAGGAGAAAATGTCCACTGGACC
43 GAGCTTTCTTTTTTGCATATTAAACTTCTGTTCCAACC
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Direct Submission
Submitted (30-APR-1998) Veterinary Microbiology, Tuskegee University, Carver Research Foundation Old Montgomery Road Tuskegee, AL 36088, USA
                                                                                                                                                                                                                                                                                                                                                                                                             AXI59660
Sequence 2988 from Patent
AXI59660
                                                                                                                                                                                                                                                    Shimkets.R.A. and Leach,M.
Nucleic acids containing single nucleotide
methods of use thereof
Patent: WO 0140521-A 2988 07-JUN-2001;
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Mammalia; Eutheria;
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Mammalia; Eutheria;
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                                                      Conservative
                                                                                                                        /note="2 of 2 allelic variants
Accession number cg42708452"
9 c 12 g 10 t
                                                                                                                                                                   /organism="Homo sapiens"
/db_xref="taxon:9606"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /Cell_type="marrow"/
/tissue_type="peritoneum"
/note="adherent peritoneal exudate cells; associated with
/note="adherent peritoneal exudate cells; associated with
                                                                                                                                                                                                                           Location/Qualifiers
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/cell_type="macrophage"
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/strain="C.D2 Vil-6"
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Primates;
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                                                     Score 21.4; DB 6;
Pred. No. 9.7e+05;
D; Mismatches 11
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Catarrhini;
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Sciurognathi; Muridae; Murinae; Mus
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AX158732
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Sequence :
AX158732
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31; Conserv
  Homo
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Location/Qualifiers
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Patent: WO 0140521-A 2060 07-JUN-2001;
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Nucleic acids containing single nucleotide
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               HSADD2S16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A method for prognosing cancer and the proteins involved Patent: WO 0206327-A 33 24-JAN-2002;
Northwestern University (US)
Location/Qualifiers
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Sequence 33 from Patent WO0206327
AX449255
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                                                                                                   AAGGAGAATTGCTGGCTCATTTTCAGGAGCCCTGCCCTTCCT 2319
                                                                            AAGGACAATGTCCTGCGAAAAATCAGGAGGCCTCCACTTCCT
D2S16 59 bp DNA sapiens beta-adducin (ADD2) gene,
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                                                                                                                               Conservative
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                                                                                                                                                                                                                                                          /organism="Homo sapiens"
/db_xref="taxon:9606"
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/db_xref="taxon:32630"
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Pred. No. 1.1e+06;
0; Mismatches 13;
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Pred. No. 9.9e+05;
0; Mismatches 16
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1 TTGCAAGGAGTGTGGRAAGGCCTTCAGCCAG
                                                                                                                                                                                        Cargill,M., Ireland,J.S. and Lander,E.S. Human single nucleotide polymorphisms Patent: WO 0166800-A 1402 13-SEP-2001; WHITEHEAD INSTITUTE FOR BIOMEDICAL RESEARCH
                                                                                                                                                                                                                                                                                                                                          31 bp | Sequence 1402 from Patent WO0166800 AX249323
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Gilligan, D.M., Lozovatsky, L. and Silberfein, A.
Direct Submission
Submitted (17-APR-1997) Internal Medicine - Section of Hematology,
Yale University School of Medicine, 333 Cedar Street, New Haven, CT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gilligan,D.M., Lozovatsky,L. and Silberfein,A. Organization of the human beta-adducin gene (ADD2) Genomics 43 (2), 141-148 (1997) 97386581
                                                    24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              32; Conserv
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Mammalia; Eutheria;
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              TTGCAGGGAGTGTGAGCAAGGCTTTAGCCAG 1388
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nilarity 64.0%;
Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /organism="Homo sapiens"
/db_xref="taxon:9606"
22. .>59
                                                                                                                   /organism="Homo sapiens"
/db_xref="taxon:9606"
6 c 11 g
                                                                                                                                                                            Location/Qualifiers
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                                                                 0.8%;
77.4%;
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Primates;
                                                   Score 21; DB Pred. No. 1.2e 1; Mismatches
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Pred. No. 1.1e+06;
0; Mismatches 18;
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                                                                 DB 6;
1.2e+06;
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RESULT 34 AX382526/c

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AR107677/C
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ORIGIN
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                                Sequence 30
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Sequence 19 from Patent WO9844121.
A84824
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synthetic
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Mehtali,M. and Legrand,V.
Mobilified Abenoviral Fiber and Targer
Patent: WO 9844121-A 19 08-OCT-1998;
MEHTALI MAJID (FR); LEGRAND VALERIE (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Zinc finger domain recognition code and uses thereof patent: WO 0208286-A 56 31-JAN-2002; Syngenta Participations AG (CH)
Location/Qualifiers
          Unknown.
                                                                  AR107677
                                                                                                                                                                                                                                                                                                                                                                                  Human adenovirus type 5.
Human adenovirus type 5
Viruses; dsDNA viruses, no
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                                                                                                                                                                                                                                                      /organism="Human adenovirus type
/Strain="ADENOVIRUS 5 (AD5)"
/isolate="OLIGONUCLEOTIDE DE SYNT
SGTV EN DKLT)"
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/db_xref="taxon:32630"
/note="PCR primer."
/note="PCR primer."
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9 c 12 g
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Pred. No. 1.3e+06;
Pred. No. 1.3e+106;
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Pred. No. 1.3e+06;
Pred. No. 5;
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WO0208286
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s 6110665.
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                                                                  PAT 14-FEB-2001
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AR165671
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ORIGIN
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AR168446/c
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Sequence 72
AR168446
                                                                              Unclassified.

Unclassified.

1 (bases 1 to 59)
Parma,D.H., Hicke,B., Bridonneau,P. and High affinity nucleic acid ligands to Patent: US 6280932-A 236 28-AUG-2001;
Location/Qualifiers
                                                                                                                                                                                                    AR165671
Sequence 236 :
AR165671
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Fenger, C.K., Granstrom, D.E., Gajadhar, A.A. and Dubey, J.P.
Sarcocystis neuronadiagnostic primer and its use in methods
equine protozoal myeloencephalitis diagnosis
Patent: US 6110665-A 30 29-AUG-2000;
 . Similarity 30; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                             1 (bases 1 to 52).

Spurr.N.K., Gray, I.C. and Stewart, L.M. Diagnosis of susceptibility to cancer Patent: US 6287854-A 72 11-SEP-2001;
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                                                    organism="unknown"
16 c 15 g
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7 c 13 g
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8 c 19 g
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Pred. No. 1.3e+06;
0; Mismatches 5;
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TRYPTASE INHIBITOR
Patent: WO 9503333-A 11 02-FEB-1995;
UCP GEN PHARNA AG (CH)
Other publication CA 2164221 19950202
Other publication AU 7532494 19950220
Other publication NO 960325 19960124
Other publication FI 96033 19960124
Other publication JP 9500532T 19970121.
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Other publication NZ 270194 960528
Other publication ZA 9500070 950707
Other publication JP 8041098 960213
                           Similarity
25; Conserv
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Sequence 7 from Patent EP0662514
A45522
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                                                                                                                                                                                                             Patent: EP 0662514-A 7 12-JUL-1995;
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Fritz, P.D., Sommerhoff, D.C. and Heim, D.J.
                                                                                                                                                                                                                                                                                          unidentified
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25; Conserv
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Sequence 11 from Patent WO9503333
A42976
                                                                                                                                                                                                                             Hiruda
                                                                                                                                                                                                                                     Hirustasin, an antistasin type serine proteinase inhibitor from
                                                                                                                                                                                                                                                                              unclassified
CAGCAGGATCCATTCTGCTTTAGTGGCAAAGC 628
                                                                                                                                                                                                                                                                                                                                                                                                                                      CAGCAAGATCTCTTCTTTAGCAGCAATGC 48
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                                                                                       /organism="unidentified"
/db_xref="taxon:32644"
13 c 7 g 1
                                                                                                                                             Location/Qualifiers
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13 c 7 g
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                                      0.8%;
78.1%;
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                                    Score 20.8;
Pred. No. 1
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Pred. No. 1.5e+06;
0; Mismatches 7;
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                                   .5e+06;
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                                                 Length 48;
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KEYWORDS
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JOURNAL
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                                                                                           RESULT 43
AX165344/c
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AUTHORS
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ORIGIN
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ORGANISM
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AR082131
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AR030794
LOCUS
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JOURNAL
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AR030794
Homo sapiens
Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                  Sequence 539 from Patent
AX165344
AX165344.1 GI:14546173
                                                                                                                                                                                                                                                                                                                                                                 AR082131
Sequence 11 from patent
AR082131
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              l Similarity
25; Conserv
                                human
                                                                                   AX165344
                                                                                                                                                                                                                                                                       Patent: US 5972698-A 11 26-OCT-1999;
                                                                                                                                                                                                                                                                                Tryptase inhibitor
                                                                                                                                                                                                                                                                                                                         Unknown.
                                                                                                                                                                                                                                                                                                                                                       AR082131.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Antistasin type serine protease inh Patent: US 5861377-A 7 19-JAN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Unknown
                                                                                                                                                                                                                                                                                                                                 Unknown.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Fritz, H., Sommerhoff, C. and Heim, J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Unclassified
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13 c 7 g
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13 c 7 g
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 Chordata; Craniata; Vertebrata; Euteleostomi; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                          Score 20.8; DB 6; Pred. No. 1.5e+06; 7;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0;
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Pred. No. 1.5e+06;
0; Mismatches 7;
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WO0138586.
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5861377.
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REFERENCE
AUTHORS
TITLE
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AX162154/c
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ORIGIN
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AUTHORS
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AX162153/c
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                                                                                                                                                                                                                             Local
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Homo sapiens
Eukaryota; Metazoa; (
Mammalia; Eutheria; H
1 (bases 1 to 51)
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                                                                                                                                                                   51
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Shimkets,R.A. and Leach,M.
Nucleic acids containing single nucleotide
                                                                                 Sequence 5482
AX162154
                                                                                                                                                                                                                L Similarity
29; Conser
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AX162153
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                                                                                                          AX162154
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                                                human..
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                                                                                                                                                                               TCTCCAGTCAGCAAGCTCTCAGCCAACATGTGTGGCCTGAGTCA 517
                                                                                                                                                                  TCTGGAGGCAGTAATCTCTCAACTCCCGGGTGTTGCTCAGGCA 9
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                                                                                                                                                                                                               0.7%;
nilarity 67.4%;
Conservative
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Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                            of use thereof
                                                                                                                                                                                                                                                                                                                                                                              WO 0140521-A 5481 07-JUN-2001;
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Accession number cg44002106"
14 C . 14 g 11 t
                                                                                                                                                                                                                                                                                                          /organism="Homo sapiens"
/db_xref="taxon:9606"
26
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Accession number cg41022625"
14 c 15 g 17 t
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/db_xref="taxon:9606"
26
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                                                                       GI:14543485
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             Chordata; Craniata; Vertebrata; Euteleostomi; Primates; Catarrhini; Hominidae; Homo.
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9: Pred. No. 1.7e-
0; Mismatches
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. No. 1.7e+06;
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Catarrhini;
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i; Hominidae;
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AR165679
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Query Match
Best Local S
Matches 26
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Best Local
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Sequence
AR165679
 I (bases 1 to 60)

I (bases 1 to 60)

Bafico,A., Varesco,L., De Benedetti,L., Caligo,M.A., Gismondi,V.Sciallero,S., Aste,H., Ferrara,G.B. and Bevilacqua,G.

Genomic PCR-SSCP analysis of the metastasis associated NM23-H1

(NME1) gene: a study on colorectal cancer

Anticancer Res. 13 (6A), 2149-2154 (1993)
                                                                                                                                                                                                                       L05010
                                                                                                                                                                                                                                                       HUMNMELE 60 bp DNA linear PRI 15-APR-HOMO sapiens (clones: ICRF c105H0563, ICRF c105E1088; subclone: BS23.13) nucleoside diphosphate kinase A chain (NME1) gene
                                                                                                                                                                                                                                                                                                                                                                                  4
                                                                                                                                                                                                                                                                                                                                                                                                                                                     ch 0.7%; Similarity 74.3%; 26; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 (bases 1 to 52)
Parma,D.H., Hicke,B., Bridonneau,P. and Gold,L.
High affinity nucleic acid ligands to lectins
Patent: US 6280932-A 244 28-AUG-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 29;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Nucleic acids containing single nucleotide methods of use thereof Patent: WO 0140521-A 5482 07-JUN-2001; Curagen Corporation (US)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Shimkets, R.A. and Leach, M.
Anticancer
94127836
                                                                                                                                    Eukaryota; Metazoa;
                                                                                                                                                       Homo sapiens
                                                                                                                                                                        Homo sapiens DNA.
                                                                                                                                                                                      nucleoside diphosphate
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                                                                                                                   Mammalia; Eutheria;
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                                                                                                                                                                                                                                            tragment.
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/db_xref="taxon:9606"
26
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /note="2 of 2 allelic variants
Accession number cg44002106"
a· 15 c 13 g 11 t
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15 c 15 g
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67.48;
                                                                                                                 Chordata;
Primates;
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Pred. No. 1.7e+06;
0; Mismatches 9;
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Pred. No. 1.7e+06;
0; Mismatches 14
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US
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                                                                                                                 Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.
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                                                                               Caligo, M.A., Gismondi, V.,
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ACCESSION
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AX158272/c
LOCUS
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ORIGIN
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Best Local S
Matches 35
JOURNAL
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Best Local Similarity 65.2%;
Matches 30; Conservative
                                    TITLE
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 Genomics
96374826
       Cloning and characterization of the murine GPI anchor gene Pigf, a homologue of the human PIGF gene Genomics 34 (3), 340-346 (1996)
                                               Kinoshita, T.
                                                          Ohishi, K., Kurimoto, Y.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AX158272
Sequence 1600 from Pate
AX158272
AX158272.1 GI:14539603
                                                                                  Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                            Mus musculus
                                                                                                                       Mus musculus DNA.
                                                                                                                                    Pig-f.
                                                                                                                                               D50270.1 GI:1753184
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                                                                                                                                                                      Mouse Pig-f DNA, exon
                                                                                                                                                                                  D50270
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Nucletc acids containing single nucleotide polymorphisms
methods of use thereof
Patent: WO 0140521-A 1600 07-JUN-2001;
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35; Conser
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/db_xref="taxon:9606"
26
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Accession number cg30793374"
14 c 14 g 13 t
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/db_xref="taxon:9606"
/map="17q22"
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                                                                                  Chordata;
Rodentia;
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Pred. No. 1.8e-
0; Mismatches
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Pred. No. 2e+06;
0; Mismatches 16;
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                                                          Inoue, N.,
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                                                                                Craniata; Vertebrata;
Sciurognathi; Muridae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Catarrhini; Hominidae;
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1.8e+06;
hes 24;
                                                          Endo, Y.,
                                                                                                                                                                                  DNA
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                                                                                                                                                                     sequence
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                                                                                                                                                                                                                                                                                                                  Length 51;
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                                                          Takeda,J.
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                                                                                              Euteleostomi;
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                                                                                  Murinae;
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                                  synthesis
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FEATURES
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ORIGIN
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REFERENCE
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Search completed: June Job time : 7203 secs
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ORGANISM
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VERSION
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AR040811/c
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AUTHORS
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                                                           2212 TGGAGACATTCTGTGTGTGATTATGCATGAGACTGTACTGG
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                                                                                                                                                                                                                                                                                                 AR040811
Sequence 40
AR040811
AR040811.1
                                                                                                                                                                                                                                                                                                                                                                                                                                             30;
                                                                                                                                                                                              1 (bases 1 to 50)
Farr, S.B. and Todd, M.D.
Methods and kits, for eukaryotic gene profiling
Patent: US 5811231-A 40 22-SEP-1998;
                                                                                                                                                                                                                                                              Unknown.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Submitted (12-APR-1995) Kazuhito Ohishi, Research Institute Microbial Diseases, Osaka University, Immunoregulation; 3-1 Yamada-oka, Suita, Osaka 565, Japan (Tel:06-875-5233, Fax:06-875-5233)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ohishi,K.
Direct Submission
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Cloning and characterization of the GPI-anchor synthesis Pig-f, murine homologue to human PIG-F gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2 (bases 1 to 52)
Ohishi, K.
                                                                                                                                                                                                                                                  Unclassified
                                                                                                                                                                                                                                                                           Unknown.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Unpublished
                                                                                                       Similarity
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30; Conservative
                                             TGCAGACATACTGAGTGTCACCGTTGAAGAGAGTGGAGTGG
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                                                                                                                                                     /organism="unknown"
16 c 8 g
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           2003,
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                                                                                          Score 20.2; DB 6;
Pred. No. 2.3e+06;
0; Mismatches 13;
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